

TONESTE STORY

FIGURE 1A

Cellular expression of $\beta Arr2-\beta gal\Delta\omega$ fusion protein in C2 clones (measured by anti- β gal ELISA)

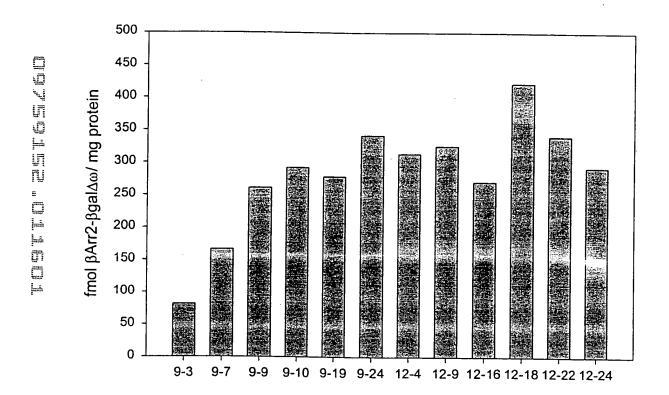


FIGURE 1B

Agonist Stimulated cAMP Response in C2 Cells Expressing β 2AR- β gal $\Delta\alpha$

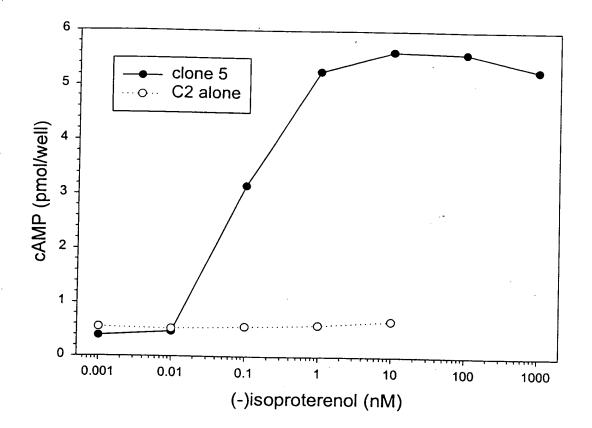
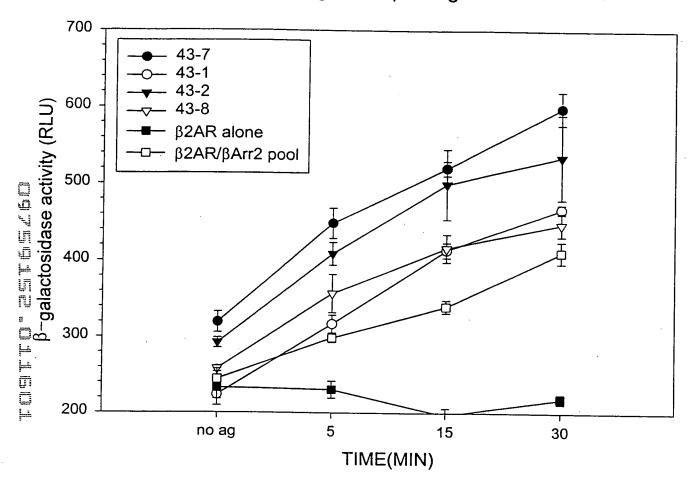


FIGURE 2

 $\beta-galactosidase$ Complementation as a Measurement for $\beta2AR-\beta gal\Delta\alpha$ interacting with $\beta Arrestin2-\beta gal\Delta\omega$ upon agonist Stimulation



 $\beta-$ galactosidase Complementation as a Measurement for $\beta 2AR-\beta gal\Delta\alpha$ Interaction with $\beta Arrestin1-\beta gal\Delta\omega$ upon Agonist Stimulation

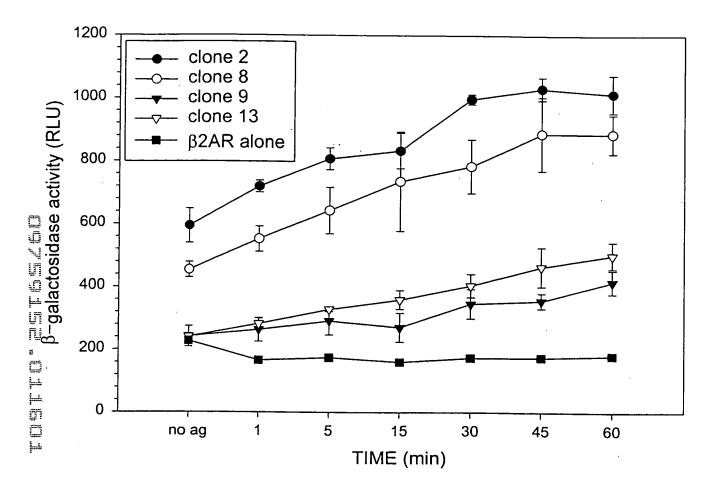


FIGURE 3B

$\beta-$ galactosidase Activity in Response to Agonist in C2 Cells Coexpressing $\beta 2AR-\beta gal\Delta\alpha$ and $\beta Arrestin2-\beta gal\Delta\omega$ Fusion Proteins

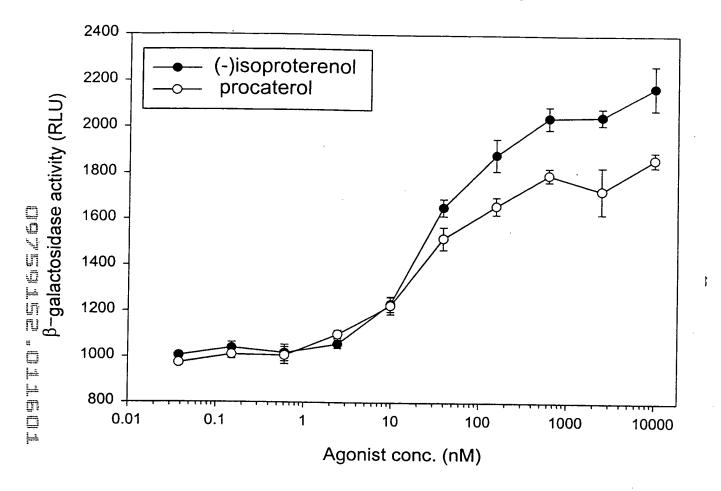


FIGURE 4A

$\beta-$ galactosidase Activity in Response to Agonist in C2 Cells Coexpressing $\beta2AR-\beta gal\Delta\alpha$ and $\beta Arrestin1-\beta gal\Delta\omega$ Fusion Proteins

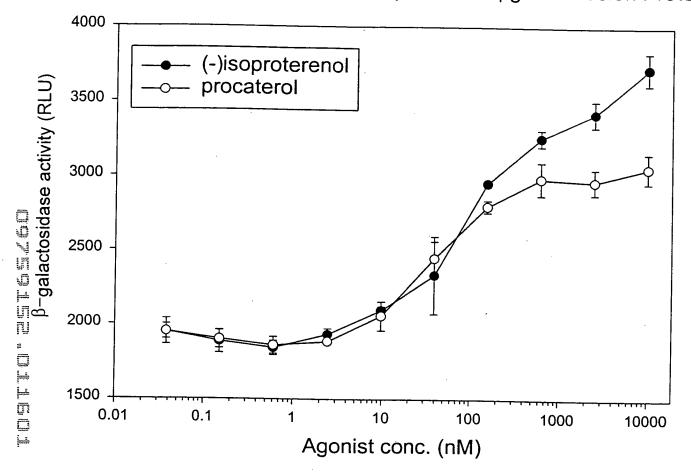


FIGURE 4B

Inhibition of β –galactosidase activity in C2 Cells Coexpressing $\beta 2AR-\beta gal\Delta\alpha$ and $\beta Arrestin2-\beta gal\Delta\omega$ Fusion Proteins

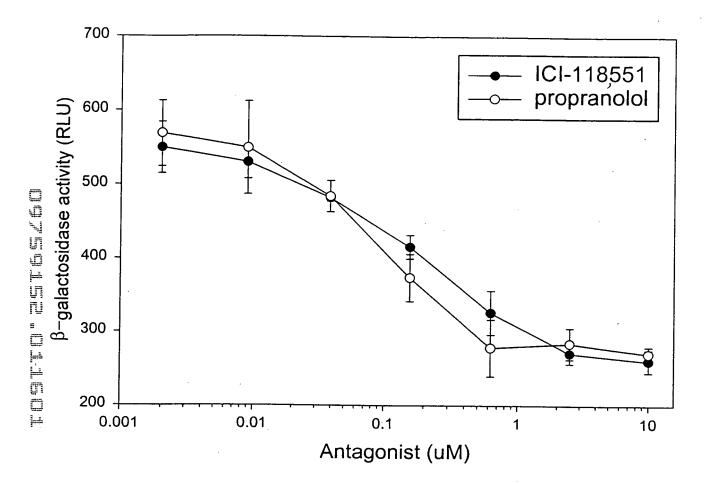
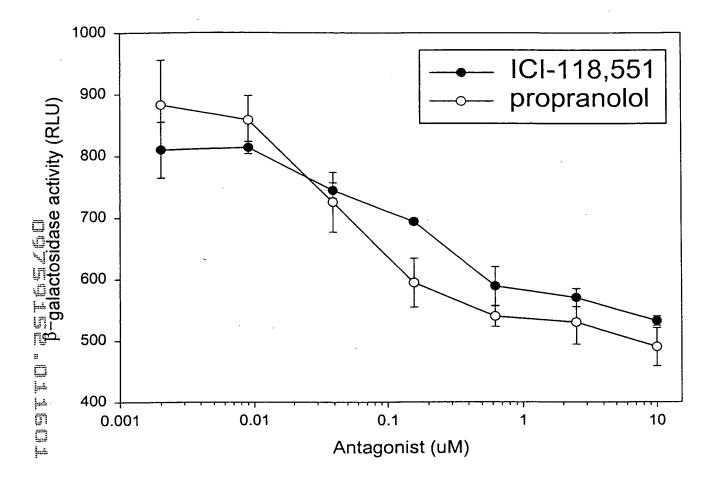


FIGURE 5A



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Figure 5B

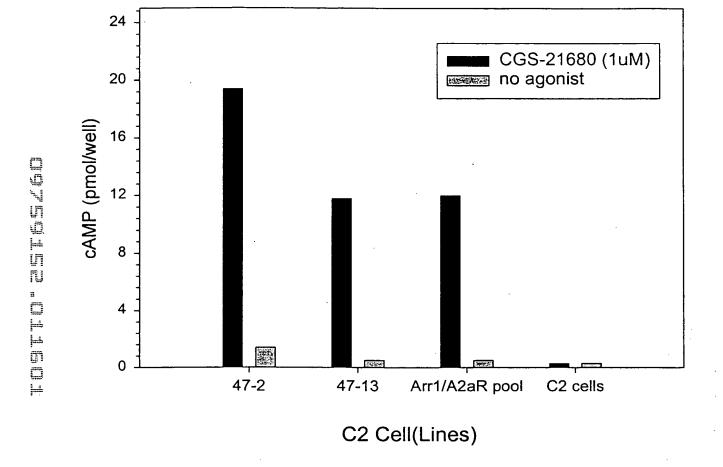


FIGURE 6

Agonist Stimulated cAMP Response in Clones or Pools of C2 Cells Expressing D1- β gal $\Delta\alpha$ and β Arrestin2- β gal $\Delta\omega$ Fusion Proteins

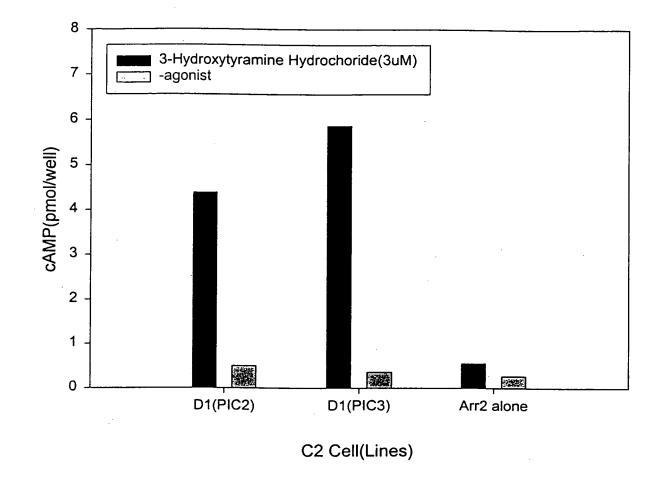


FIGURE 7

$β_2$ AR-βgalΔω and βarr2-βgalΔα Interaction in HEK293 Clones in Response to Isoproterenol Treatment (1 μM)

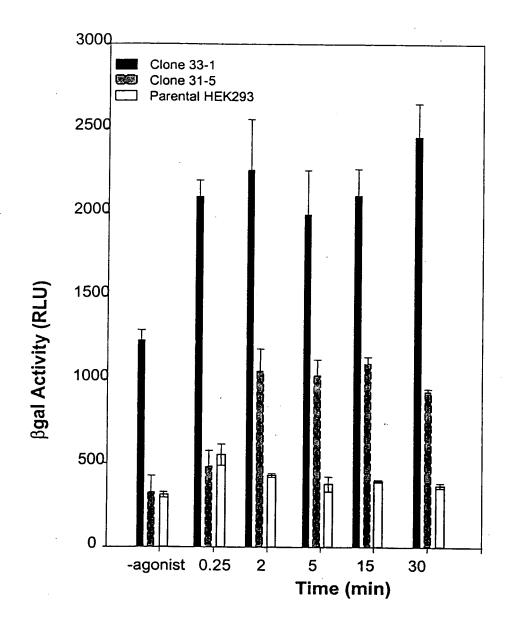
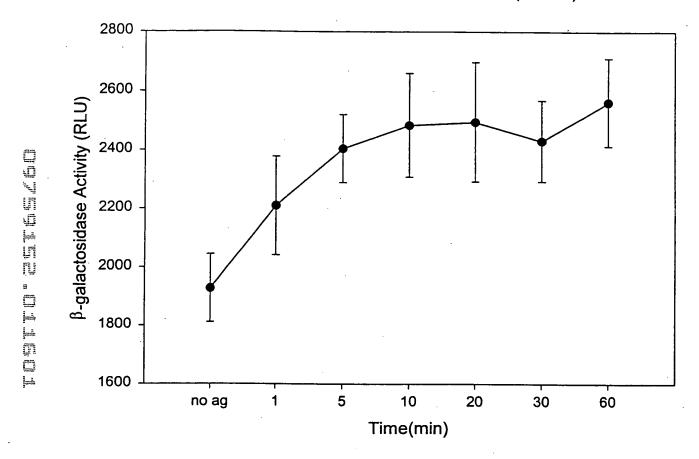
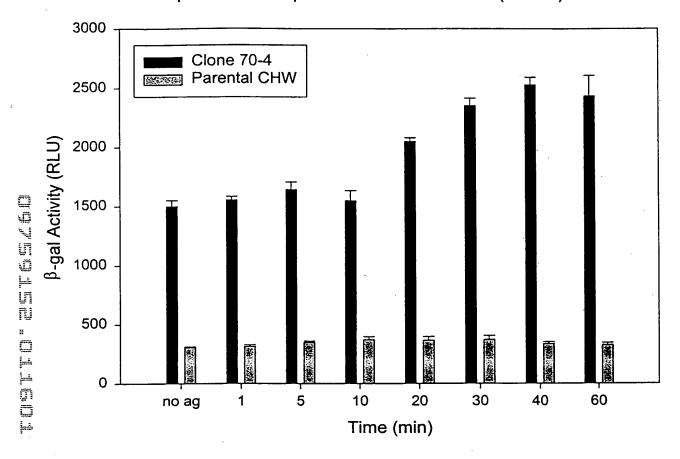


FIGURE 8A

β2AR-βgalΔα and βArr1-βgalΔω Interaction in a CHO Pool in Response to Isoproterenol Treatment(10uM)



$\beta 2AR-\beta gal\Delta\alpha$ and $\beta Arr2-\beta gal\Delta\omega$ Interaction in CHW Clone in Response to Isoproterenol Treatment (10uM)



 β –galactosidase Complementation as a Measurement for Adrenergic Receptor Homodimerization in HEK 293 Cells Coexpressing β 2AR- β gal Δ α and β 2AR- β gal Δ ω.

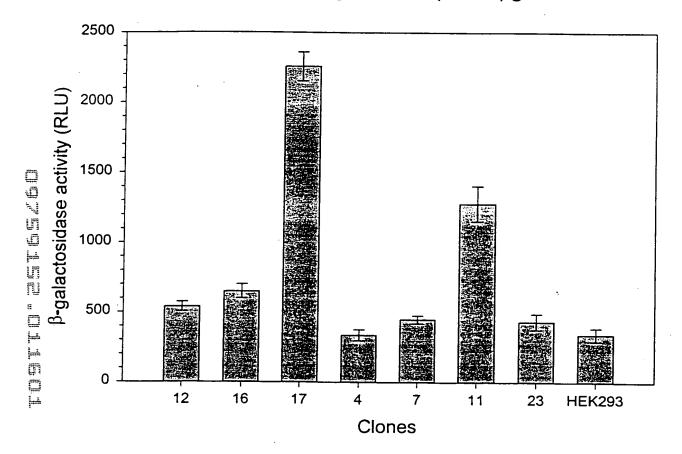
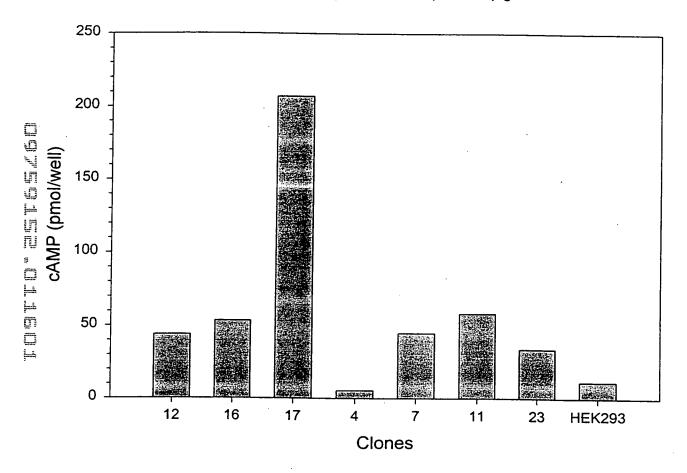


FIGURE 9A

Agonist Stimulated cAMP Response in HEK 293 Cells Coexpressing $\beta 2AR-\beta gal\Delta\alpha$ and $\beta 2AR-\beta gal\Delta\omega$



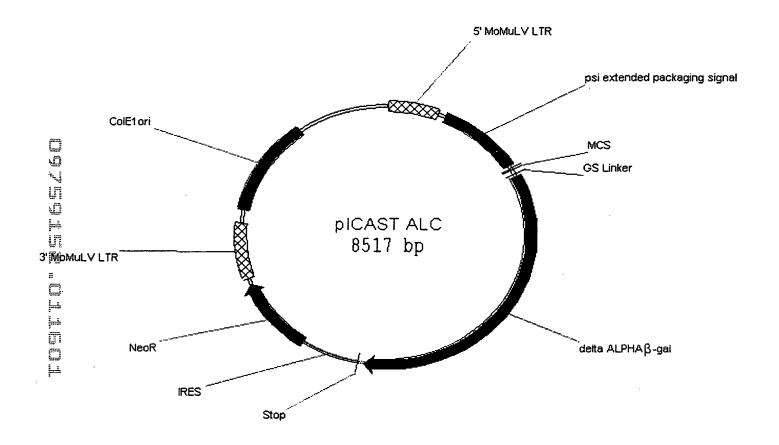


Figure 10A

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51	CCCCGGCTCA GGGGCCGAGT					
101	GGATATCTGT CCTATAGACA					
151	GGTCCCCAGA CCAGGGGTCT			TTTCTAGAGA AAAGATCTCT		
201	GTTTCCAGGG CAAAGGTCCC					·
251	TAACCAATCA ATTGGTTAGT					
301 301	GCTCAATAAA CGAGTTATTT			CGGGGCGCCA GCCCCGCGGT		
				AATAAACCCT TTATTTGGGA		
3 1553	CATCCGACTT			GAGGGTCTCC CTCCCAGAGG		
45 1	TGACTACCCG ACTGATGGGC			GGGGGCTCGT CCCCCGAGCA		
501 51	GAGACCCCTG CTCTGGGGAC			CACCGGGAGG GTGGCCCTCC		
551				AGTGTCTATG TCACAGATAC	ACTGATTTTA TGACTAAAAT	
601	TGCGCCTGCG ACGCGGACGC			AGCTCTGTAT TCGAGACATA		·
651	CGTGGTGGAA GCACCACCTT	CTGACGAGTT GACTGCTCAA	CTGAACACCC GACTTGTGGG	GGCCGCAACC CCGGCGTTGG	CTGGGAGACG GACCCTCTGC	
701		AAACCCCCGG	CAAAAACACC	GGGCTGGACT	CCTTCCCTCA	
	CGATGTGGAA GCTACACCTT	TCCGACCCCG AGGCTGGGGC	TCAGGATATG AGTCCTATAC	TGGTTCTGGT ACCAAGACCA	AGGAGACGAG TCCTCTGCTC	
	AACCTAAAAC TTGGATTTTG	AGTTCCCGCC TCAAGGGCGG	TCCGTCTGAA AGGCAGACTT	TTTTTGCTTT AAAAACGAAA	CGGTTTGGAA GCCAAACCTT	
851	CCGAAGCCGC GGCTTCGGCG	GCGTCTTGTC CGCAGAACAG	TGCTGCAGCA ACGACGTCGT	TCGTTCTGTG AGCAAGACAC	TTGTCTCTGT AACAGAGACA	
	CTGACTGTGT GACTGACACA	TTCTGTATTT AAGACATAAA	GTCTGAAAAT CAGACTTTTA	TAGGGCCAGA ATCCCGGTCT	CTGTTACCAC GACAATGGTG	

95 L	TOCCTTANGT			AGATGTCGAG TCTACAGCTC		
	MOGGAATIOA	AMCIGORATO	CATIONCOIL	ICIACAGCIC		
1001	ACAACCAGTC	GGTAGATGTC	AAGAAGAGAC	GTTGGGTTAC	CTTCTGCTCT	
1001				CAACCCAATG		·
1051	GCAGAATGGC	CAACCTTTAA	CGTCGGATGG	CCGCGAGACG	GCACCTTTAA	
	CGTCTTACCG	GTTGGAAATT	GCAGCCTACC	GGCGCTCTGC	CGTGGAAATT	
1101	CCGAGACCTC					
	GGCTCTGGAG	TAGTGGGTCC	AATTCTAGTT	CCAGAAAAGT	GGACCGGGCG	
1151	ATGGACACCC			ACTGGACCT		
	TACCIGIGGG	1CTGGTCCAG	GGGAIGIAGC	ACIGGACCCI	TCGGAACCGA	
1201	TTTGACCCCC	СТСССТСССТ	СВВССССТТТ	GTACACCCTA	ACCUTCUCC	
1201				CATGTGGGAT		•
			. 			
1251	TCCTCTTCCT	CCATCCGCCC	CGTCTCTCCC	CCTTGAACCT	CCTCGTTCGA	·
1				GGAACTTGGA		•
<u></u>						
13701	CCCCGCCTCG	ATCCTCCCTT	TATCCAGCCC	TCACTCCTTC	TCTAGGCGCC	•
ū		TAGGAGGGAA	ATAGGTCGGG	AGTGAGGAAG	AGATCCGCGG	
#						
1951				ATAGGGCGAT		
Ñ	CCGGCGAGAT	CGGGTAATTA	TGCTGAGTGA	TATCCCGCTA	AGCTTAGTCC	
 a						
1401	CCTTGGCGCG					·
Ħ	GGAACCGCGC	GGCCTAGGAA	TTAATTCGCG	TTAACCCTCC	ACCGCCATCG	•
+2						
T +2		VIT		A V V A		
	CTCGAGATGG					
				CGGCAGCACC		
+2	R P S	Q Q L F	R S L N	G E W	R F A	
1.501		CARCACHUAC	CCACCCECAA	mccccn a mcc	CCCMMMCCCM	
1501	TCGCCCTTCC			ACCGCTTACC		
+2	WFPA	PEA	V P E S	S W T. F.	C D To	
1551	GGTTTCCGGC					
	CCAAAGGCCG	TGGTCTTCGC	CACGGCCTTT	CGACCGACCT	CACGCTAGAA	
+2	PEAG	YVT	V P S	N M O N	1 H G Y	
1601	CCTGAGGCCG	 ътъстстсст	CCTCCCCTCA	AACTGGCAGA	TCCACCCTTA	
1001				TTGACCGTCT		
·						
+2	DAP	IYTN	VTY	РІТ	V N P	
1651	CGATGCGCCC					•
	GCTACGCGGG	TAGATGTGGT	TGCACTGGAT	AGGGTAATGC	CAGTTAGGCG	

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+2 P F V P T E N P T G C Y S L T F N
1701 CGTTTGTTCC CACGGAGAAT CCGACGGGTT GTTACTCGCT CACATTTAAT GCAAACAAGG GTGCCTCTTA GGCTGCCCAA CAATGAGCGA GTGTAAATTA
+2 V D E S W L Q E G Q T R I F D G
1751 GTTGATGAAA GCTGGCTACA GGAAGGCCAG ACGCGAATTA TTTTTGATGG CAACTACTTT CGACCGATGT CCTTCCGGTC TGCGCTTAAT AAAAACTACC
+2 V N S A F H L W C N G R W V G Y
1801 CGTTAACTCG GCGTTTCATC TGTGGTGCAA CGGGCGCTGG GTCGGTTACG GCAATTGAGC CGCAAAGTAG ACACCACGTT GCCCGCGACC CAGCCAATGC
+2 G Q D S R L P S E F D L S A F L R
CGGTCCTGTC AGCAAACGGC AGACTTAAAC TGGACTCGCG TAAAAATGCG +2 A G E N R L A V M V L R W S D G S
+2 A G E N R L A V M V L R W S D G S
1901 GCCGGAGAAA ACCGCCTCGC GGTGATGGTG CTGCGCTGGA GTGACGGCAG CGGCCTCTTT TGGCGGAGCG CCACTACCAC GACGCGACCT CACTGCCGTC
+2 Y L E D Q D M W R M S G I F R D
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+2 V S L L H K P T T Q I S D F H V A
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+2 T R F N D D F S R A V L E A E V Q
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12 M C C B I D B V I D V M V C I W
+2 M C G E L R D Y L R V T V S L W
2101 GATGTGCGGC GAGTTGCGTG ACTACCTACG GGTAACAGTT TCTTTATGGC CTACACGCCG CTCAACGCAC TGATGGATGC CCATTGTCAA AGAAATACCG
+2 Q G E T Q V A S G T A P F G G E I
2151 AGGGTGAAAC GCAGGTCGCC AGCGGCACCG CGCCTTTCGG CGGTGAAATT
TCCCACTTTG CGTCCAGCGG TCGCCGTGGC GCGGAAAGCC GCCACTTTAA
+2 I D E R G G Y A D R V T L R L N V
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+2 E N P K L W S A E I P N L Y R A
2251 CGAAAACCCG AAACTGTGGA GCGCCGAAAT CCCGAATCTC TATCGTGCGG
GCTTTTGGGC TTTGACACCT CGCGGCTTTA GGGCTTAGAG ATAGCACGCC

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+2 V V E L H T A D G T L I E A E A C 2301 TGGTTGAACT GCACACCGCC GACGGCACGC TGATTGAAGC AGAAGCCTGC ACCAACTTGA CGTGTGGCGG CTGCCGTGCG ACTAACTTCG TCTTCGGACG +2 D V G F R E V R I E N G L I I I N	
+2 D V G F R E V R T F N C * * * * * * * * * * * * * * * * * *	
+2 D V G F R E V R I E N G L L L N	
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+2 G K P L L I R G V N R H E H H P	
2401 CGGCAAGCCG TTGCTGATTC GAGGCGTTAA CCGTCACGAG CATCATCCTC GCCGTTCGGC AACGACTAAG CTCCGCAATT GGCAGTGCTC GTAGTAGGAG	
+2 L H G Q V M D E Q T M V Q D I L L	 -
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F	
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+2 E A N I E T H G M V P M N R L T D	
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+2 T D I I C P M Y A R V D E D Q P	
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+2	2 E T R P L I L C E Y A H A M G N S	
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+2	2 LGGFAKY WQAFRQYPR	
_	TCTTGGCGGT TTCGCTAAAT ACTGGCAGGC GTTTCGTCAG TATCCCCGTT AGAACCGCCA AAGCGATTTA TGACCGTCCG CAAAGCAGTC ATAGGGGCAA	
		·
+2	2 L Q G G F V W D W V D Q S L I K Y	
3051	TACAGGGCGG CTTCGTCTGG GACTGGGTGG ATCAGTCGCT GATTAAATAT ATGTCCCGCC GAAGCAGACC CTGACCCACC TAGTCAGCGA CTAATTTATA	
	DENG NPW SAY GGDF GDT	* * *
3 10 1 亞	GATGAAAACG GCAACCCGTG GTCGGCTTAC GGCGGTGATT TTGGCGATAC CTACTTTTGC CGTTGGGCAC CAGCCGAATG CCGCCACTAA AACCGCTATG	
异 切+2	P N D R Q F C M N G L V F A D R	
3 15 1		
- <u>-</u>	TPHPALTEAKH QQQ FFQ	
32 9 1 =	CGCCGCATCC AGCGCTGACG GAAGCAAAAC ACCAGCAGCA GTTTTTCCAG GCGGCGTAGG TCGCGACTGC CTTCGTTTTG TGGTCGTCGT CAAAAAGGTC	
+2	FRLSGQTIEVTSEYLFR	
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+2	H S D N E L L H W M V A L D G K	
	TCATAGCGAT AACGAGCTCC TGCACTGGAT GGTGGCGCTG GATGGTAAGC AGTATCGCTA TTGCTCGAGG ACGTGACCTA CCACCGCGAC CTACCATTCG	·
 +2	PLASGEV PLDV APQ GKQ	
	CGCTGGCAAG CGGTGAAGTG CCTCTGGATG TCGCTCCACA AGGTAAACAG	
+2	LIEL PEL PQPESAG QLW	
3401	TTGATTGAAC TGCCTGAACT ACCGCAGCCG GAGAGCGCCG GGCAACTCTG AACTAACTTG ACGGACTTGA TGGCGTCGGC CTCTCGCGGC CCGTTGAGAC	
+2	L T V R V V Q P N A T A W S E A	
3451	GCTCACAGTA CGCGTAGTGC AACCGAACGC GACCGCATGG TCAGAAGCCG CGAGTGTCAT GCGCATCACG TTGGCTTGCG CTGGCGTACC AGTCTTCGGC	

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+2 T L P A A S H A I P H L T T S E M
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+2 S G F L S Q M W I G D K K Q L L T
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F=
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+2 A A G H Y Q A E A A L L Q C T A D
3801 CGGCGGGCCA TTACCAGGCC GAAGCAGCGT TGTTGCAGTG CACGGCAGAT GCCGCCCGGT AATGGTCCGG CTTCGTCGCA ACAACGTCAC GTGCCGTCTA
+2 T L A D A V L I T T A H A W Q H Q
3851 ACACTTGCTG ATGCGGTGCT GATTACGACC GCTCACGCGT GGCAGCATCA TGTGAACGAC TACGCCACGA CTAATGCTGG CGAGTGCGCA CCGTCGTAGT
+2 G K T L F I S R K T Y R I D G S
3901 GGGGAAAACC TTATTTATCA GCCGGAAAAC CTACCGGATT GATGGTAGTG CCCCTTTTGG AATAAATAGT CGGCCTTTTG GATGGCCTAA CTACCATCAC
+2 G Q M A I T V D V E V A S D T P H
3951 GTCAAATGGC GATTACCGTT GATGTTGAAG TGGCGAGCGA TACACCGCAT CAGTTTACCG CTAATGGCAA CTACAACTTC ACCGCTCGCT ATGTGGCGTA
+2 P A R I G L N C Q L A Q V A E R V
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+2 N W L G L G P Q E N Y P D R L T
4051 AAACTGGCTC GGATTAGGGC CGCAAGAAAA CTATCCCGAC CGCCTTACTG TTTGACCGAG CCTAATCCCG GCGTTCTTTT GATAGGGCTG GCGGAATGAC

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+2 A A C F D R W D L P L S D M Y T P	
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+2 Y V F P S E N G L R C G T R E L N	
4151 TACGTCTTCC CGAGCGAAAA CGGTCTGCGC TGCGGGACGC GCGAATTGAA ATGCAGAAGG GCTCGCTTTT GCCAGACGCG ACGCCCTGCG CGCTTAACTT	
+2 Y G P H Q W R G D F Q F N I S R	
4201 TTATGGCCCA CACCAGTGGC GCGGCGACTT CCAGTTCAAC ATCAGCCGCT AATACCGGGT GTGGTCACCG CGCCGCTGAA GGTCAAGTTG TAGTCGGCGA	
	
+2 Y S Q Q L M E T S H R H L L H A	
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+2 E E G T W L N I D G F H M G I G G	
4301 GAAGAAGCA CATGGCTGAA TATCGACGGT TTCCATATGG GGATTGGTGG CTTCTTCCGT GTACCGACTT ATAGCTGCCA AAGGTATACC CCTAACCACC	
+2 D D S W S P S V S A E F Q L S A	
7	
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+2 G R Y H Y Q L V W C Q K R S D Y K	
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+2 D E D L D H H H H H R	
4451 GATGAGGACC TCGACCATCA TCATCATCAT CACCGGTAAT AATAGGTAGA	
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4501 TAAGTGACTG ATTAGATGCA TTGATCCCTC GACCAATTCC GGTTATTTTC ATTCACTGAC TAATCTACGT AACTAGGGAG CTGGTTAAGG CCAATAAAAG	
4551 CACCATATTG CCGTCTTTTG GCAATGTGAG GGCCCGGAAA CCTGGCCCTG	
GTGGTATAAC GGCAGAAAAC CGTTACACTC CCGGGCCTTT GGACCGGGAC	
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4651 CAAGGTCTGT TGAATGTCGT GAAGGAAGCA GTTCCTCTGG AAGCTTCTTG	
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4701 AAGACAAACA ACGTCTGTAG CGACCCTTTG CAGGCAGCGG AACCCCCCAC TTCTGTTTGT TGCAGACATC GCTGGGAAAC GTCCGTCGCC TTGGGGGGGTG	
4751 CTGGCGACAG GTGCCTCTGC GGCCAAAAGC CACGTGTATA AGATACACCT	
GACCGCTGTC CACGGAGACG CCGGTTTTCG GTGCACATAT TCTATGTGGA	

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4801		CACAACCCCA GTGTTGGGGT			
4851	AAGAGTCAAA TTCTCAGTTT	TGGCTCTCCT ACCGAGAGGA			
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4951		GTGTTTAGTC CACAAATCAG			
5001	CACGGGGACG GTGCCCCTGC	TGGTTTTCCT ACCAAAAGGA			
5051	AACAAGATGG TTGTTCTACC	ATTGCACGCA TAACGTGCGT			
5 fg1	AAGCCGATAC	ACTGGGCACA TGACCCGTGT	TGTCTGTTAG	CCGACGAGAC	TACGGCGGCA
5 15 1	GTTCCGGCTG CAAGGCCGAC	TCAGCGCAGG AGTCGCGTCC	GGCGCCCGGT CCGCGGGCCA	TCTTTTTGTC AGAAAAACAG	AAGACCGACC TTCTGGCTGG
52 0 1 TU		CCTGAATGAA GGACTTACTT		TCCGTCGCGC	
52 <u>5</u> 1		CGGGCGTTCC GCCCGCAAGG			·
53 <u>0</u> 1		GACTGGCTGC CTGACCGACG			•
5351 		CCTTGCTCCT GGAACGAGGA			
5401	ATGCGGCGGC TACGCCGCCG	TGCATACGCT ACGTATGCGA			
5451	AGCGAAACAT TCGCTTTGTA	CGCATCGAGC GCGTAGCTCG			
5501	TCGATCAGGA AGCTAGTCCT	TGATCTGGAC ACTAGACCTG	GAAGAGCATC CTTCTCGTAG	AGGGGCTCGC TCCCCGAGCG	GCCAGCCGAA CGGTCGGCTT
5551	CTGTTCGCCA GACAAGCGGT	GGCTCAAGGC CCGAGTTCCG			
5601	GACCCATGGC CTGGGTACCG	GATGCCTGCT CTACGGACGA			
5651		CATCGACTGT GTAGCTGACA	CCGGCCGACC		
5701	GACATAGCGT CTGTATCGCA	TGGCTACCCG ACCGATGGGC			

5751	GGCTGACCGC CCGACTGGCG	TTCCTCGTGC AAGGAGCACG	TTTACGGTAT AAATGCCATA	CGCCGCTCCC	GATTCGCAGC CTAAGCGTCG	
5801		CTATCGCCTT GATAGCGGAA	GAACTGCTCA	TCTTCTGAGC AGAAGACTCG	GGGACTCTGG CCCTGAGACC	
5851	GGTTCGCATC CCAAGCGTAG	GATAAAATAA CTATTTTATT	AAGATTTTAT TTCTAAAATA	TTAGTCTCCA AATCAGAGGT	CTTTTTCCCC	
5901	GGAATGAAAG CCTTACTTTC	ACCCCACCTG TGGGGTGGAC	TAGGTTTGGC ATCCAAACCG	AAGCTAGCTT TTCGATCGAA	AAGTAACGCC TTCATTGCGG	
5951	ATTTTGCAAG TAAAACGTTC	GCATGGAAAA CGTACCTTTT	ATACATAACT TATGTATTGA	GAGAATAGAG CTCTTATCTC	TTCAAGTCTA	
6001	CAAGGTCAGG GTTCCAGTCC		ACAGCTGAAT TGTCGACTTA			
6 0 51	GTGGTAAGCA CACCATTCGT		CGGCTCAGGG GCCGAGTCCC			
Ł	TGAATATGGG ACTTATACCC	GGTTTGTCCT	TATCTGTGGT ATAGACACCA	AAGCAGTTCC TTCGTCAAGG	ACGGGGCCGA	**
6 1 751	CAGGGCCAAG GTCCCGGTTC				AGTCGTCAAA	
6 2 01	CTAGAGAACC GATCTCTTGG	ATCAGATGTT TAGTCTACAA	TCCAGGGTGC AGGTCCCACG	CCCAAGGACC GGGTTCCTGG	TGAAATGACC ACTTTACTGG	
6251	CTGTGCCTTA GACACGGAAT	TTTGAACTAA AAACTTGATT	CCAATCAGTT GGTTAGTCAA	CGCTTCTCGC GCGAAGAGCG	TTCTGTTCGC AAGACAAGCG	
6501 	GCGCTTCTGC CGCGAAGACG	TCCCCGAGCT AGGGGCTCGA	CAATAAAAGA GTTATTTTCT	GCCCACAACC CGGGTGTTGG	CCTCACTCGG GGAGTGAGCC	
6351	GGCGCCAGTC CCGCGGTCAG					
		CGTCAACGTA	GGCTGAACAC	CAGAGCGACA	AGGAACCCTC	
	GGTCTCCTCT CCAGAGGAGA	CTCACTAACT	GATGGGCAGT	CGCCCCAGA	AAGTAAGTAC	
	CAGCATGTAT GTCGTACATA	GTTTTAATTA 	AACCAAAAA	AAGAATTCAT	AAATGTAATT	
	ATGGCCATAG TACCGGTATC	AACGTAATTA 	CTTAGCCGGT	TGCGCGCCCC	TCTCCGCCAA	
	TGCGTATTGG ACGCATAACC	GCGAGAAGGC 	GAAGGAGCGA	GTGACTGAGC	GACGCGAGCC	
6651	TCGTTCGGCT AGCAAGCCGA	GCGGCGAGCG CGCCGCTCGC	GTATCAGCTC CATAGTCGAG	ACTCAAAGGC TGAGTTTCCG	GGTAATACGG CCATTATGCC	

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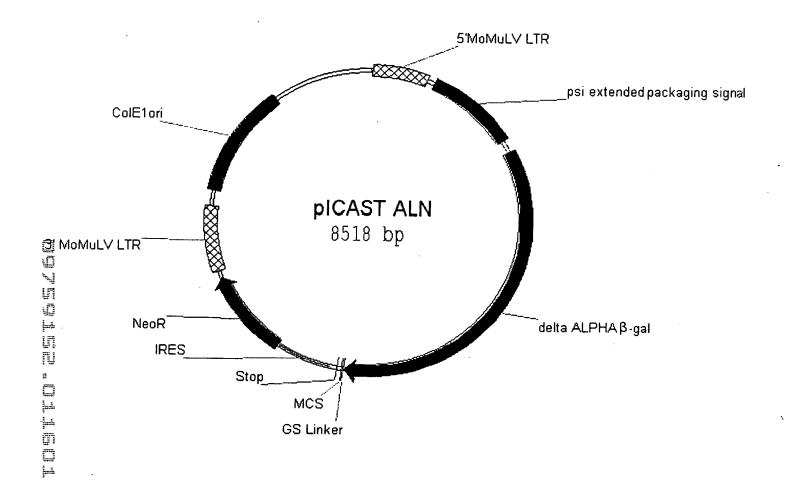


Figure 11A

1	CTGCAGCCTG GACGTCGGAC	AATATGGGCC TTATACCCGG	AAACAGGATA TTTGTCCTAT	TCTGTGGTAA AGACACCATT	GCAGTTCCTG CGTCAAGGAC	
51	CCCCGGCTCA GGGGCCGAGT	GGGCCAAGAA CCCGGTTCTT	CAGATGGAAC GTCTACCTTG	AGCTGAATAT TCGACTTATA	GGGCCAAACA CCCGGTTTGT	
101	GGATATCTGT CCTATAGACA	GGTAAGCAGT CCATTCGTCA	TCCTGCCCG	GCTCAGGGCC CGAGTCCCGG	AAGAACAGAT TTCTTGTCTA	
151	GGTCCCCAGA CCAGGGGTCT	TGCGGTCCAG ACGCCAGGTC	CCCTCAGCAG GGGAGTCGTC	TTTCTAGAGA AAAGATCTCT	TGGTAGTCTA	
201	GTTTCCAGGG CAAAGGTCCC	TGCCCCAAGG ACGGGGTTCC	ACCTGAAATG TGGACTTTAC	ACCCTGTGCC TGGGACACGG	TTATTTGAAC AATAAACTTG	
251	TAACCAATCA ATTGGTTAGT	GTTCGCTTCT CAAGCGAAGA	CGCTTCTGTT GCGAAGACAA	CGCGCGCTTC GCGCGCGAAG	TGCTCCCCGA	
301 		AGAGCCCACA TCTCGGGTGT	ACCCCTCACT TGGGGAGTGA	CGGGGCGCCA GCCCCGCGGT	GTCCTCCGAT CAGGAGGCTA	
351 迈 止	TGACTGAGTC	GCCCGGGTAC CGGGCCCATG	CCGTGTATCC	AATAAACCCT TTATTTGGGA	GAACGTCAAC	
401 TU	CATCCGACTT GTAGGCTGAA	GTGGTCTCGC CACCAGAGCG	TGTTCCTTGG ACAAGGAACC	GAGGGTCTCC CTCCCAGAGG	TCTGAGTGAT AGACTCACTA	
45 1 □	TGACTACCCG ACTGATGGGC	TCAGCGGGGG AGTCGCCCCC	TCTTTCATTT AGAAAGTAAA	GGGGGCTCGT CCCCCGAGCA	CCGGGATCGG GGCCCTAGCC	
501 (T) -(D) -	GAGACCCCTG CTCTGGGGAC	CCCAGGGACC GGGTCCCTGG	ACCGACCCAC TGGCTGGGTG	CACCGGGAGG GTGGCCCTCC	CAAGCTGGCC	
551	AGCAACTTAT TCGTTGAATA	CTGTGTCTGT GACACAGACA	CCGATTGTCT GGCTAACAGA	AGTGTCTATG TCACAGATAC	ACTGATTTTA TGACTAAAAT	
601	TGCGCCTGCG ACGCGGACGC	TCGGTACTAG AGCCATGATC	TTAGCTAACT AATCGATTGA	AGCTCTGTAT TCGAGACATA	CTGGCGGACC GACCGCCTGG	
651	CGTGGTGGAA GCACCACCTT	CTGACGAGTT GACTGCTCAA	CTGAACACCC GACTTGTGGG	GGCCGCAACC CCGGCGTTGG	CTGGGAGACG GACCCTCTGC	
701	TCCCAGGGAC AGGGTCCCTG	AAACCCCCGG (CAAAAACACC	GGGCTGGACT	CCTTCCCTCA	
751	CGATGTGGAA GCTACACCTT	TCCGACCCCG '	TCAGGATATG AGTCCTATAC	TGGTTCTGGT ACCAAGACCA	AGGAGACGAG TCCTCTGCTC	
801	AACCTAAAAC TTGGATTTTG	AGTTCCCGCC T	TCCGTCTGAA AGGCAGACTT	TTTTTGCTTT AAAAACGAAA	CGGTTTGGAA GCCAAACCTT	
851	CCGAAGCCGC (GGCTTCGGCG (GCGTCTTGTC 1	TGCTGCAGCA	TCGTTCTGTG	ттстстстст	
901	CTGACTGTGT GACTGACACA	AAGACATAAA (CAGACTTTTA	ATCCCGGTCT	GACAATGGTG	

951					CGGCTCGCTC GCCGAGCGAG
1001	ACAACCAGTC TGTTGGTCAG				CTTCTGCTCT GAAGACGAGA
1051	GCAGAATGGC CGTCTTACCG				GCACCTTTAA CGTGGAAATT
1101	CCGAGACCTC GGCTCTGGAG				CCTGGCCGC GGACCGGGCG
1151	ATGGACACCC TACCTGTGGG				AGCCTTGGCT TCGGAACCGA
1201					AGCCTCCGCC TCGGAGGCGG
1251 11					CCTCGTTCGA GGAGCAAGCT
1301 	CCCCGCCTCG GGGGCGGAGC				TCTAGGCGCC AGATCCGCGG
ş	GGCCGCTCTA CCGGCGAGAT	GCCCATTAAT CGGGTAATTA	ACGACTCACT TGCTGAGTGA	ATAGGGCGAT TATCCCGCTA	TCGAACACCA AGCTTGTGGT
1401					CCTCGAGATG GGAGCTCTAC
1451 					ATCGCCCTTC TAGCGGGAAG
1501	CCAACAGTTA GGTTGTCAAT				TGGTTTCCGG ACCAAAGGCC
1551	CACCAGAAGC GTGGTCTTCG				TCCTGAGGCC AGGACTCCGG
1601	GATACTGTCG CTATGACAGC				ACGATGCGCC TGCTACGCGG
1651			TAGGGTAATG	CCAGTTAGGC	GGCAAACAAG
1701					TGTTGATGAA ACAACTACTT
1751					GCGTTAACTC CGCAATTGAG
1801					GGCCAGGACA CCGGTCCTGT
1851					CGCCGGAGAA GCGGCCTCTT

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1901				AGTGACGGCA TCACTGCCGT		
1951	AGATCAGGAT TCTAGTCCTA			TTTCCGTGAC AAAGGCACTG		
2001	TGCATAAACC ACGTATTTGG			TCCATGTTGC AGGTACAACG		
2051	AATGATGATT TTACTACTAA			GCTGAAGTTC CGACTTCAAG		
2101	CGAGTTGCGT GCTCAACGCA			TTCTTTATGG AAGAAATACC		
2151	CGCAGGTCGC GCGTCCAGCG			GCGGTGAAAT CGCCACTTTA		
2 20 1				CGTCTGAACG GCAGACTTGC	AGCTTTTGGG	
2 25 1	GAAACTGTGG CTTTGACACC	TCGCGGCTTT			GTGGTTGAAC CACCAACTTG	
14 2 30 1 Tu		CGACGGCACG		CAGAAGCCTG GTCTTCGGAC	CGATGTCGGT GCTACAGCCA	·
2351				CTGCTGCTGA GACGACGACT	ACGGCAAGCC	
2401	GTTGCTGATT CAACGACTAA			GCATCATCCT CGTAGTAGGA		
	AGGTCATGGA TCCAGTACCT			ATATCCTGCT TATAGGACGA		
2501	AACAACTTTA TTGTTGAAAT			TATCCGAACC ATAGGCTTGG		
2551	GTACACGCTG CATGTGCGAC			TGTGGTGGAT ACACCACCTA		
		GCCGTACCAC	GGTTACTTAG	CAGACTGGCT	ACTAGGCGCG	
2651		GCTACTCGCT	TGCGCATTGC	GCTTACCACG	TCGCGCTAGC	
2701	TAATCACCCG ATTAGTGGGC			GGGGAATGAA CCCCTTACTT		
2751	GCGCTAATCA CGCGATTAGT			TCAAATCTGT AGTTTAGACA		
2801	CGCCCGGTGC GCGGGCCACG			GACACCACGG CTGTGGTGCC		

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2851		ATGTACGCGC TACATGCGCG				
2901	TGCCGAAATG ACGGCTTTAC	GTCCATCAAA CAGGTAGTTT				
2951	CCGCTGATCC GGCGACTAGG	TTTGCGAATA AAACGCTTAT				
3001	TTTCGCTAAA AAAGCGATTT	TACTGGCAGG ATGACCGTCC				
3051	GCTTCGTCTG CGAAGCAGAC	GGACTGGGTG CCTGACCCAC				
3101	GGCAACCCGT CCGTTGGGCA	GGTCGGCTTA CCAGCCGAAT				
3151	TCGCCAGTTC AGCGGTCAAG	TGTATGAACG ACATACTTGC				
3 20 1 ∰	CAGCGCTGAC GTCGCGACTG	CCTTCGTTTT	GTGGTCGTCG		CAAGGCAAAT	
3 25 1		CCATCGAAGT GGTAGCTTCA	GACCAGCGAA CTGGTCGCTT	TACCTGTTCC ATGGACAAGG	GTCATAGCGA CAGTATCGCT	
3301 <u></u> -		CTGCACTGGA GACGTGACCT		-		
3 35 1 		GCCTCTGGAT CGGAGACCTA				
	CTGCCTGAAC GACGGACTTG	TACCGCAGCC ATGGCGTCGG				
3451	ACGCGTAGTG TGCGCATCAC	CAACCGAACG GTTGGCTTGC				
3501	GCGCCTGGCA CGCGGACCGT	GCAGTGGCGT CGTCACCGCA				
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3601	CATCGAGCTG GTAGCTCGAC	GGTAATAAGC CCATTATTCG				****
3651		GTGGATTGGC CACCTAACCG				
3701		GGGCACGTGG	CGACCTATTG	CTGTAACCGC	ATTCACTTCG	
3751	GACCCGCATT CTGGGCGTAA	GACCCTAACG CTGGGATTGC				

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3801	ATTACCAGGC TAATGGTCCG	CGAAGCAGCG GCTTCGTCGC	TTGTTGCAGT AACAACGTCA	GCACGGCAGA CGTGCCGTCT	TACACTTGCT ATGTGAACGA	
3851	GATGCGGTGC CTACGCCACG	TGATTACGAC ACTAATGCTG	CGCTCACGCG GCGAGTGCGC	TGGCAGCATC ACCGTCGTAG	AGGGGAAAAC TCCCCTTTTG	
3901	CTTATTTATC GAATAAATAG	AGCCGGAAAA TCGGCCTTTT	CCTACCGGAT GGATGGCCTA	TGATGGTAGT ACTACCATCA	GGTCAAATGG CCAGTTTACC	
3951	CGATTACCGT GCTAATGGCA	TGATGTTGAA ACTACAACTT	GTGGCGAGCG CACCGCTCGC	ATACACCGCA TATGTGGCGT	TCCGGCGCGG AGGCCGCGCC	
4001	ATTGGCCTGA 'TAACCGGACT	ACTGCCAGCT TGACGGTCGA	GGCGCAGGTA CCGCGTCCAT	GCAGAGCGGG CGTCTCGCCC	TAAACTGGCT ATTTGACCGA	
4051	CGGATTAGGG GCCTAATCCC	CCGCAAGAAA GGCGTTCTTT	ACTATCCCGA TGATAGGGCT	CCGCCTTACT GGCGGAATGA	GCCGCCTGTT CGGCGGACAA	
4피1 빌 닠	TTGACCGCTG AACTGGCGAC	GGATCTGCCA CCTAGACGGT	TTGTCAGACA AACAGTCTGT	TGTATACCCC ACATATGGGG	GTACGTCTTC CATGCAGAAG	
4 5 1 ©	CCGAGCGAAA GGCTCGCTTT	ACGGTCTGCG TGCCAGACGC	CTGCGGGACG GACGCCCTGC	CGCGAATTGA GCGCTTAACT	ATTATGGCCC TAATACCGGG	
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4251 	TCGTTGACTA	CCTTTGGTCG	GTAGCGGTAG	ACGACGTGCG	CCTTCTTCCG	
43 0 1 (1) -(2)	ACATGGCTGA TGTACCGACT	ATATCGACGG TATAGCTGCC	TTTCCATATG AAAGGTATAC	GGGATTGGTG CCCTAACCAC	GCGACGACTC CGCTGCTGAG	
4351	CTGGAGCCCG GACCTCGGGC	TCAGTATCGG AGTCATAGCC	CGGAATTCCA GCCTTAAGGT	GCTGAGCGCC CGACTCGCGG	GGTCGCTACC CCAGCGATGG	
	ATTACCAGTT TAATGGTCAA	CCAGACCACA	GTTTTTTCTA	GACCTCCACC	ACCGTCGTCC	
4451	CCTTGGCGCG GGAACCGCGC	CCGGATCCTT GGCCTAGGAA	AATTAACAAT TTAATTGTTA	TGACCGGTAA ACTGGCCATT	TAATAGGTAG ATTATCCATC	
	ATAAGTGACT TATTCACTGA	CTAATCTACG	TAACTAGGGA	GCTGGTTAAG	GCCAATAAAA	
		CGGCAGAAAA	CCGTTACACT	CCCGGGCCTT	TGGACCGGGA	
		GCTCGTAAGG	ATCCCCAGAA	AGGGGAGAGC	GGTTTCCTTA	
	GCAAGGTCTG CGTTCCAGAC	AACTTACAGC	ACTTCCTTCG	TCAAGGAGAC	CTTCGAAGAA	
4701	GAAGACAAAC CTTCTGTTTG	TTGCAGACAT	CGCTGGGAAA	CGTCCGTCGC	CTTGGGGGGT	

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4751	CCTGGCGACA GGACCGCTGT				AAGATACACC TTCTATGTGG
4801	TGCAAAGGCG ACGTTTCCGC				ATAGTTGTGG TATCAACACC
4851	AAAGAGTCAA TTTCTCAGTT				GCTGAAGGAT CGACTTCCTA
4901	GCCCAGAAGG CGGGTCTTCC				CTCGGTGCAC GAGCCACGTG
4951	ATGCTTTACA TACGAAATGT				GCCCCCGAA CGGGGGGCTT
5001					TACCATGATT ATGGTACTAA
	GAACAAGATG CTTGTTCTAC		TCCAAGAGGC	CGGCGAACCC	TGGAGAGGCT ACCTCTCCGA
5 10 1 11 -	ATTCGGCTAT				GATGCCGCCG CTACGGCGGC
5 15 1					CAAGACCGAC GTTCTGGCTG
5201	GACAGGCCAC				GGCTATCGTG CCGATAGCAC
-	GCTGGCCACG				GTTGTCACTG CAACAGTGAC
5301		GGACTGGCTG CCTGACCGAC			GCAGGATCTC CGTCCTAGAG
5351	CTGTCATCTC GACAGTAGAG	ACCTTGCTCC TGGAACGAGG			
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5551	ACTGTTCGCC	AGGCTCAAGG	CGCGCATGCC GCGCGTACGG	CGACGGCGAG	GATCTCGTCG CTAGAGCAGC
5601		GCTACGGACG	TTGCCGAATA AACGGCTTAT	TCATGGTGGA AGTACCACCT	
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5701	GGACATAGCG	TTGGCTACCC	GTGATATTGC	TGAAGAGCTT	GGCGGCGAAT	•
	CCTGTATCGC	AACCGATGGG	CACTATAACG	ACTTCTCGAA	CCGCCGCTTA	
5751	GGGCTGACCG	C##CC#CC#C	CTTTACCCTA	中ででででででですべて	CCATTCCCAC	
2/31				AGCGGCGAGG		
	CCCGACTGGC	GAAGGAGCAC	GAAATGCCAT	AGCGGCGAGG	GCTAAGCGTC	
5801	CGCATCGCCT					
	GCGTAGCGGA	AGATAGCGGA	AGAACTGCTC	AAGAAGACTC	GCCCTGAGAC	
5851	GGGTTCGCAT	CGATAAAATA	AAAGATTTTA	TTTAGTCTCC	AGAAAAAGGG	
				AAATCAGAGG		
E 0.0.1	CCCAAMCAAA	CACCCACCT	CTACCTTTCC	C	ጥአ አ ርጥአ አ ሮርር	
3901	GGGAATGAAA			GTTCGATCGA		
	CCCTTACTT	CIGGGGIGGA	CATCCAAACC	GIICGAICGA	ATTCATTGCG	
5951	CATTTTGCAA	GGCATGGAAA	AATACATAAC	TGAGAATAGA	GAAGTTCAGA	
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6 d d 1	TCAAGGTCAG	GAACAGATGG	AACAGCTGAA	TATGGGCCAA	ACAGGATATC	
<u>u</u>	AGTTCCAGTC	CTTGTCTACC	TTGTCGACTT	ATACCCGGTT	TGTCCTATAG	
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6051	ጥርጥርርጥ ን አርር	NGTTCCTCCC	CCGCCTCAGG	GCCAAGAACA	CATCCAACAC	
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	CTGAATATGG					
N	GACTTATACC	CGGTTTGTCC	TATAGACACC	ATTCGTCAAG	GACGGGGCCG	
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L	AGTCCCGGTT	CTTGTCTACC	AGGGGTCTAC	GCCAGGTCGG	GAGTCGTCAA	
6201	TCTAGAGAAC	CATCAGATGT	TTCCAGGGTG	CCCCAAGGAC	CTGAAATGAC	
TI.				GGGGTTCCTG		
I						
6 के के 1	CCTGTGCCTT	δ ΨΨΨΩ δ Δ CΨδ	ACCAATCAGT	тесеттете	СТТСТСТТСС	
0231				AGCGAAGAGC		•
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6401	TAAACCCTCT	TGCAGTTGCA	TCCGACTTGT	GGTCTCGCTG	TTCCTTGGGA	
	•			CCAGAGCGAC		
6451	GGGTCTCCTC	тсастсаттс	ACTACCCGTC	AGCGGGGGTC	ጥጥጥርልጥጥርልጥ	
0431					AAAGTAAGTA	
	DADDADGAG	ACICACIAAC				
6501	GCAGCATGTA					
					TAAATGTAAT	
6551	AATGGCCATA					
	TTACCGGTAT	CAACGTAATT	ACTTAGCCGG	TTGCGCGCCC	CTCTCCGCCA	•
6601	TTGCGTATTG	GCGCTCTTCC	GCTTCCTCGC	TCACTGACTC	GCTGCGCTCG	
					CGACGCGAGC	

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6701	GTTATCCACA CAATAGGTGT	GAATCAGGGG CTTAGTCCCC				
 6751	GCCAGCAAAA CGGTCGTTTT	GGCCAGGAAC CCGGTCCTTG				
6801	CATAGGCTCC GTATCCGAGG	GCCCCCTGA CGGGGGGACT				
6851	GAGGTGGCGA CTCCACCGCT	AACCCGACAG TTGGGCTGTC				
6901	GAAGCTCCCT CTTCGAGGGA	CGTGCGCTCT GCACGCGAGA				
J==						
6951 N	CTGTCCGCCT GACAGGCGGA	TTCTCCCTTC AAGAGGGAAG				
7 0 01 Ū	CTGTAGGTAT GACATCCATA	CTCAGTTCGG GAGTCAAGCC				
7051 TU 5		CCCCGTTCAG GGGGCAAGTC				
	CGTCTTGAGT GCAGAACTCA	CCAACCCGGT GGTTGGGCCA				
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7151 □	CACTGGTAAC GTGACCATTG	AGGATTAGCA TCCTAATCGT				
7201		GTGGCCTAAC CACCGGATTG				
7251	ATCTGCGCTC TAGACGCGAG	TGCTGAAGCC ACGACTTCGG				
7301	TTGATCCGGC AACTAGGCCG	AAACAAACCA TTTGTTTGGT			AAACAAACGT	
7351	AGCAGCAGAT TCGTCGTCTA	TACGCGCAGA ATGCGCGTCT		-	TCCTTTGATC	

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7455						
		TCTAATAGTT	TTTCCTAGAA	GTGGATCTAG	GAAAACGCCG	
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7551	ATGCTTAATC	AGTGAGGCAC	CTATCTCAGC	GATCTGTCTA		

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7601	CCATAGTTGC GGTATCAACG	CTGACTCCCC	GTCGTGTAGA CAGCACATCT	TAACTACGAT ATTGATGCTA	ACGGGAGGGC TGCCCTCCCG	
7651	TTACCATCTG AATGGTAGAC	GCCCCAGTGC CGGGGTCACG	TGCAATGATA ACGTTACTAT	CCGCGAGACC GGCGCTCTGG	CACGCTCACC GTGCGAGTGG	
7701	GGCTCCAGAT CCGAGGTCTA		TAAACCAGCC ATTTGGTCGG		CGGCTCGCGT	
7751	GAAGTGGTCC CTTCACCAGG	TGCAACTTTA ACGTTGAAAT	TCCGCCTCCA AGGCGGAGGT	TCCAGTCTAT AGGTCAGATA	TAATTGTTGC	
7801	CGGGAAGCTA GCCCTTCGAT	GAGTAAGTAG CTCATTCATC	TTCGCCAGTT AAGCGGTCAA	AATAGTTTGC TTATCAAACG	GCAACGTTGT CGTTGCAACA	
7851	TGCCATTGCT ACGGTAACGA	ACAGGCATCG TGTCCGTAGC	TGGTGTCACG ACCACAGTGC	CTCGTCGTTT GAGCAGCAAA	GGTATGGCTT CCATACCGAA	
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9 9 01		CGTCACAATA	CACTCATGGT GTGAGTACCA			
8051	CTCTTACTGT GAGAATGACA				ACCACTCATG	
8 1 01 (1) (2)	TCAACCAAGT AGTTGGTTCA	CATTCTGAGA GTAAGACTCT	ATAGTGTATG TATCACATAC	CGGCGACCGA GCCGCTGGCT	GTTGCTCTTG	
8 11 51			ATACCGCGCC TATGGCGCGG			
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8251	CCGCTGTTGA GGCGACAACT					
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8451	TCTCATGAGC AGAGTACTCG	GGATACATAT CCTATGTATA	TTGAATGTAT AACTTACATA	TTAGAAAAAT AATCTTTTTA	AAACAAATAG TTTGTTTATC	
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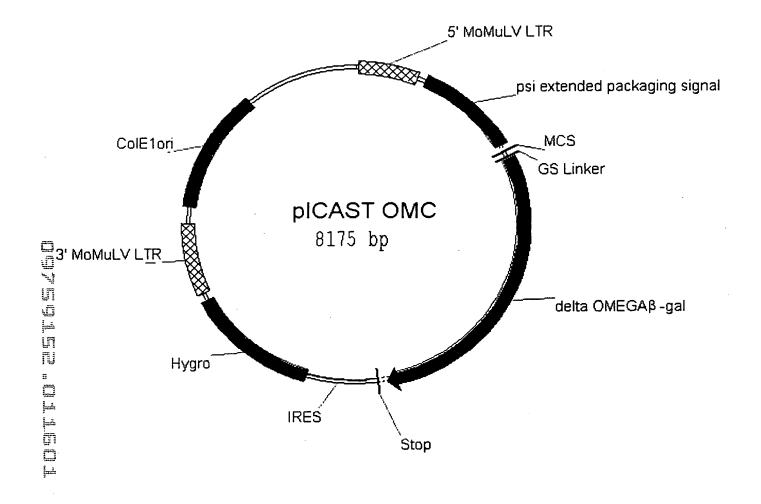


Figure 12A

1					A GCAGTTCCTG CCTCAAGGAC	
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201					C TTATTTGAAC G AATAAACTTG	
					TGCTCCCGA G ACGAGGGGCT	
301					A GTCCTCCGAT F CAGGAGGCTA	
351 W					CTTGCAGTTG A GAACGTCAAC	
#01 U				•	C TCTGAGTGAT G AGACTCACTA	
451					CCGGGATCGG A GGCCCTAGCC	_ ~
591 0	*		and the second s		G CAAGCTGGCC C GTTCGACCGG	
551					G ACTGATTTA C TGACTAAAAT	
601					CTGGCGGACC A GACCGCCTGG	
651					C CTGGGAGACG G GACCCTCTGC	
701	AGGGTCCCTG	AAACCCCCGG	CAAAAACACC	GGGCTGGACT	A GGAAGGGAGT I CCTTCCCTCA	
751	GCTACACCTT	AGGCTGGGGC	AGTCCTATAC	ACCAAGACCA	F AGGAGACGAG A TCCTCTGCTC	
801				AAAAACGAAA	T CGGTTTGGAA A GCCAAACCTT	
851					G TTGTCTCTGT C AACAGAGACA	
901	GACTGACACA	AAGACATAAA	CAGACTTTTA	ATCCCGGTCT	A CTGTTACCAC T GACAATGGTG	

951		TTGACCTTAG AACTGGAATC			
1001	ACAACCAGTC TGTTGGTCAG			_	CTTCTGCTCT GAAGACGAGA
1051	GCAGAATGGC CGTCTTACCG	CAACCTTTAA GTTGGAAATT			
1101	CCGAGACCTC GGCTCTGGAG	ATCACCCAGG TAGTGGGTCC			
1151	ATGGACACCC TACCTGTGGG				AGCCTTGGCT TCGGAACCGA
1201	TTTGACCCCC AAACTGGGGG				AGCCTCCGCC TCGGAGGCGG
1251		CCATCCGCCC GGTAGGCGGG			CCTCGTTCGA GGAGCAAGCT
13 0 1 ① 	CCCCGCCTCG GGGGCGGAGC	ATCCTCCCTT TAGGAGGGAA			
13 5]			TGCTGAGTGA		AGCTTAGTCC
1401	-	CCGGATCCTT GGCCTAGGAA			•
1451 	CTCGAGATGG GAGCTCTACC	GCGTGATTAC CGCACTAATG			
15 0 1					GCAGCACATC CGTCGTGTAG
1551	CCCCTTTCGC GGGGAAAGCG				CGATCGCCCT GCTAGCGGGA
1601	TCCCAACAGT AGGGTTGTCA			-	CCTGGTTTCC GGACCAAAGG
1651			TTTCGACCGA		CTTCCTGAGG GAAGGACTCC
1701	CCGATACTGT GGCTATGACA				TTACGATGCG AATGCTACGC
1751	CCCATCTACA GGGTAGATGT			TGCCAGTTAG	CGCCGTTTGT GCGGCAAACA
1801	TCCCACGGAG AGGGTGCCTC				AATGTTGATG TTACAACTAC
1851	AAAGCTGGCT TTTCGACCGA				TGGCGTTAAC ACCGCAATTG

1901	TCGGCGTTTC AGCCGCAAAG	ATCTGTGGTG TAGACACCAC				
1951	CAGTCGTTTG GTCAGCAAAC	CCGTCTGAAT GGCAGACTTA				
2001	AAAACCGCCT TTTTGGCGGA	CGCGGTGATG GCGCCACTAC				
2051	GAAGATCAGG CTTCTAGTCC	ATATGTGGCG TATACACCGC				
2101	GCTGCATAAA CGACGTATTT	CCGACTACAC GGCTGATGTG				
2151	TTAATGATGA AATTACTACT	TTTCAGCCGC AAAGTCGGCG				
2201		GTGACTACCT CACTGATGGA				
	AACGCAGGTC TTGCGTCCAG	GCCAGCGGCA CGGTCGCCGT				
p.	AGCGTGGTGG TCGCACCACC	TTATGCCGAT				
2351		GGAGCGCCGA CCTCGCGGCT				
2401 		GCCGACGGCA CGGCTGCCGT				
2451	CAAAGGCGCT	GGTGCGGATT CCACGCCTAA	CTTTTACCAG		CTTGCCGTTC	
2501	CCGTTGCTGA GGCAACGACT	TTCGAGGCGT AAGCTCCGCA				
2551	TCAGGTCATG AGTCCAGTAC	GATGAGCAGA CTACTCGTCT	CGATGGTGCA GCTACCACGT	GGATATCCTG CCTATAGGAC	CTGATGAAGC GACTACTTCG	,
2601	AGAACAACTT TCTTGTTGAA	TAACGCCGTG ATTGCGGCAC				
2651	TGGTACACGC ACCATGTGCG	TGTGCGACCG ACACGCTGGC				
2701	TATTGAAACC ATAACTTTGG				GATGATCCGC CTACTAGGCG	
2751		CCGCTACTCG	CTTGCGCATT	GCGCTTACCA	CGTCGCGCTA	
		GCTCACACTA	GTAGACCAGC	GACCCCTTAC	TTAGTCCGGT	

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2851		CACGACGCGC GTGCTGCGCG				
2901	CCCGCCCGGT GGGCGGGCCA	GCAGTATGAA CGTCATACTT				
2951	ATTATTTGCC TAATAAACGG	CGATGTACGC GCTACATGCG				
3001		TGGTCCATCA ACCAGGTAGT				
3051	GCCCGCTGAT CGGGCGACTA	CCTTTGCGAA GGAAACGCTT				
3101		AATACTGGCA TTATGACCGT	CCGCAAAGCA	GTCATAGGGG		
2 🚟 1	CGGCTTCGTC					
JFJ1		ACCCTGACCC				
3 20 1	ACGGCAACCC	GTGGTCGGCT	TACGGCGGTG	ATTTTGGCGA	TACGCCGAAC	
ü		CACCAGCCGA				•
<u></u>						
3251	GATCGCCAGT					1
N	CTAGCGGTCA	AGACATACTT	GCCAGACCAG	AAACGGCTGG	CGTGCGGCGT	
2201						
3301		ACGGAAGCAA TGCCTTCGTT				
<u></u>						
3351	TATCCGGGCA	AACCATCGAA	GTGACCAGCG	AATACCTGTT	CCGTCATAGC	
m		TTGGTAGCTT				
- []						
3401	GATAACGAGC					
	CTATTGCTCG			GACCTACCAT		
	AAGCGGTGAA			7C77CCT777		
2421		CACGGAGACC				• •
3501	AACTGCCTGA	ACTACCGCAG	CCGGAGAGCG	CCGGGCAACT	CTGGCTCACA	
	TTGACGGACT	TGATGGCGTC	GGCCTCTCGC	GGCCCGTTGA	GACCGAGTGT	
3551	GTACGCGTAG					
	CATGCGCATC	ACGTTGGCTT	GCGCTGGCGT	ACCAGTCTTC	GGCCCGTGTA	
3601	CAGCGCCTGG	CACCACTCCC	CTCTGGCGGA	AAACCTCACT	CTCACCCTCC	
3001		GTCGTCACCG			·	
3651	CCGCCGCGTC	CCACGCCATC	CCGCATCTGA	CCACCAGCGA	AATGGATTTT	
	GGCGGCGCAG					
3701	TGCATCGAGC					
	ACGTAGCTCG	ACCCATTATT			TCAGTCCGAA	·
3751	TCTTTCACAG			_	ACGCCCCTCC	
5,51		TACACCTAAC				•

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3801	GCGATCAGTT CGCTAGTCAA	CACCCGTGTC GTGGGCACAG	GATAGATCTG CTATCTAGAC	AACAGAAACT TTGTCTTTGA	CATTTCCGAA GTAAAGGCTT	
3851	GAAGACCTAG CTTCTGGATC	TCGACCATCA AGCTGGTAGT	TCATCATCAT AGTAGTAGTA	CACCGGTAAT GTGGCCATTA	AATAGGTAGA TTATCCATCT	
3901	TAAGTGACTG ATTCACTGAC	ATTAGATGCA TAATCTACGT	TTTCGACTAG AAAGCTGATC	ATCCCTCGAC TAGGGAGCTG	CAATTCCGGT GTTAAGGCCA	
3951	TATTTTCCAC ATAAAAGGTG	CATATTGCCG GTATAACGGC	TCTTTTGGCA AGAAAACCGT	ATGTGAGGGC TACACTCCCG	CCGGAAACCT GGCCTTTGGA	
4001	GGCCCTGTCT CCGGGACAGA	TCTTGACGAG AGAACTGCTC	CATTCCTAGG GTAAGGATCC	GGTCTTTCCC CCAGAAAGGG	GAGAGCGGTT	
4051	AGGAATGCAA TCCTTACGTT	GGTCTGTTGA CCAGACAACT				
4 <u>10</u> 1		ACAAACAACG TGTTTGTTGC	AGACATCGCT	GGGAAACGTC	CGTCGCCTTG	
4151 		CGCTGTCCAC	CCTCTGCGGC GGAGACGCCG	CAAAAGCCAC GTTTTCGGTG	CACATATTCT	
4201 TU	TACACCTGCA ATGTGGACGT				TCAACCTATC	
4 <u>2</u> 51	TTGTGGAAAG AACACCTTTC					
4301 =	AAGGATGCCC TTCCTACGGG					
	GTGCACATGC CACGTGTACG	AAATGTACAC	AAATCAGCTC	CAATTTTTTG	CAGATCCGGG	
	CCCGAACCAC GGGCTTGGTG	CCCCTGCACC	AAAAGGAAAC	TTTTTGTGCT	ACTATTATGG	
		GACTTGAGTG	GCGCTGCAGA	CAGCTCTTCA	AAGACTAGCT	
_'		TCGCAGAGGC	TGGACTACGT	CGAGAGCCTC	CCGCTTCTTA	
		GTCGAAGCTA	CATCCTCCCG	CACCTATACA	GGACGCCCAT	·
	AATAGCTGCG TTATCGACGC	GGCTACCAAA	GATGTTTCTA	GCAATACAAA	TAGCCGTGAA	
	TGCATCGGCC ACGTAGCCGG	CGCGAGGGCT	AAGGCCTTCA	CGAACTGTAA	CCCCTTAAAT	
4701	GCGAGAGCCT CGCTCTCGGA	GACCTATTGC CTGGATAACG	TAGAGGGCGG	CACGTGTCCC	ACAGTGCAAC	

4751	CAAGACCTGC GTTCTGGACG			GTTCTGCAGC CAAGACGTCG		
4801	GGCCATGGAT CCGGTACCTA			TAGCCAGACG ATCGGTCTGC		
4851	GCCCATTCGG CGGGTAAGCC			ACACTACATG TGTGATGTAC		
4901	ATATGCGCGA TATACGCGCT			CACTGGCAAA GTGACCGTTT		
4951	CGACACCGTC GCTGTGGCAG	AGTGCGTCCG TCACGCAGGC	TCGCGCAGGC AGCGCGTCCG	TCTCGATGAG AGAGCTACTC	CTGATGCTTT GACTACGAAA	
5001	GGGCCGAGGA CCCGGCTCCT			TCGTGCACGC AGCACGTGCG		
50 5 1 -[7]				ATAACAGCGG TATTGTCGCC		
51 <u>01</u>			TAAGGGTTAT		AACATCTTCT TTGTAGAAGA	
-U7 5151 -	TCTGGAGGCC	GTGGTTGGCT	TGTATGGAGC ACATACCTCG		CTACTTCGAG GATGAAGCTC	
5201					CGTATATGCT GCATATACGA	
5251 ———					GGCAATTTCG CCGTTAAAGC	
5301				ACGCAATCGT TGCGTTAGCA	CCGATCCGGA GGCTAGGCCT	
5351	GCCGGGACTG CGGCCCTGAC	TCGGGCGTAC AGCCCGCATG	ACAAATCGCC TGTTTAGCGG	CGCAGAAGCG GCGTCTTCGC	CGGCCGTCTG GCCGGCAGAC	
5401	GACCGATGGC CTGGCTACCG	TGTGTAGAAG ACACATCTTC	TACTCGCCGA ATGAGCGGCT	TAGTGGAAAC ATCACCTTTG	CGACGCCCCA GCTGCGGGGT	
		CTCCCGTTTC	CTTATCTCAT	CTACGGCTGG	GGGATCTATC CCCTAGATAG	
5501		TTCTAAAATA	AATCAGAGGT	CTTTTTCCCC	CCTTACTTTC	
5551	ACCCCACCTG TGGGGTGGAC			TTCATTGCGG	TAAAACGTTC	
5601		TATGTATTGA	CTCTTATCTC	TTCAAGTCTA	CAAGGTCAGG GTTCCAGTCC	
5651		ACAGCTGAAT	ATGGGCCAAA	CAGGATATCT		·

5701	GTTCCTGCCC CAAGGACGGG					
5751	CCAAACAGGA GGTTTGTCCT			TGCCCCGGCT ACGGGGCCGA	GTCCCGGTTC	
5801	AACAGATGGT TTGTCTACCA			TCAGCAGTTT AGTCGTCAAA		
5851	ATCAGATGTT TAGTCTACAA			TGAAATGACC ACTTTACTGG		
5901			GCGAAGAGCG	AAGACAAGCG		
5951	TCCCCGAGCT	CAATAAAAGA	GCCCACAACC			
60 0 1				TGTATCCAAT ACATAGGTTA		
<u> </u>	GCAGTTGCAT CGTCAACGTA	GGCTGAACAC	CAGAGCGACA	AGGAACCCTC	CCAGAGGAGA	
t行 6把1 。	GAGTGATTGA CTCACTAACT	CTACCCGTCA	GCGGGGGTCT		CAGCATGTAT	
6 <u>15</u> 1	CAAAATTAAT GTTTTAATTA			TTTACATTAA AAATGTAATT		
6201	AACGTAATTA	GAATCGGCCA CTTAGCCGGT	ACGCGCGGGG TGCGCGCCCC	AGAGGCGGTT TCTCCGCCAA	TGCGTATTGG ACGCATAACC	
6251	CGCTCTTCCG			CTGCGCTCGG GACGCGAGCC		
6301	GCGGCGAGCG CGCCGCTCGC			GGTAATACGG CCATTATGCC		
6351	AATCAGGGGA TTAGTCCCCT			GAGCAAAAGG CTCGTTTTCC		
6401	GCCAGGAACC CGGTCCTTGG			GCGTTTTTCC CGCAAAAAGG		
6451		CTCGTAGTGT	TTTTAGCTGC	CTCAAGTCAG GAGTTCAGTC	TCCACCGCTT	· · · · · · · · · · · · · · · · · · ·
6501	ACCCGACAGG	ACTATAAAGA	TACCAGGCGT	TTCCCCCTGG		
6551		GACAAGGCTG	GGACGGCGAA	TGGCCTATGG	ACAGGCGGAA	
6601	TCTCCCTTCG	GGAAGCGTGG	CGCTTTCTCA	TAGCTCACGC		

6651			CGCTCCAAGC GCGAGGTTCG		
6701	CCCGTTCAGC GGGCAAGTCG		CGCCTTATCC GCGGAATAGG		
6751	CAACCCGGTA GTTGGGCCAT		TATCGCCACT ATAGCGGTGA		
6801	GGATTAGCAG CCTAATCGTC		GTAGGCGGTG CATCCGCCAC		CTTGAAGTGG
6851			TAGAAGAACA ATCTTCTTGT		TCTGCGCTCT
6901	GCTGAAGCCA CGACTTCGGT	CAATGGAAGC	GAAAAAGAGT CTTTTTCTCA	ACCATCGAGA	ACTAGGCCGT
695 j	AACAAACCAC TTGTTTGGTG	CGCTGGTAGC	GGTGGTTTTT CCACCAAAAA	TTGTTTGCAA	GCAGCAGATT
ļ.	ACGCGCAGAA TGCGCGTCTT	TTTTTCCTAG	TCAAGAAGAT AGTTCTTCTA	CCTTTGATCT GGAAACTAGA	TTTCTACGGG AAAGATGCCC
-7 -	GTCTGACGCT CAGACTGCGA	CAGTGGAACG			
710	GATTATCAAA CTAATAGTTT		ACCTAGATCC TGGATCTAGG		
715	TTGCGGCCGC AACGCCGGCG		AAAGTATATA TTTCATATAT		
7201	GTTACCAATG CAATGGTTAC		GAGGCACCTA CTCCGTGGAT		CTGTCTATTT
7251	CGTTCATCCA GCAAGTAGGT		ACTCCCCGTC TGAGGGGCAG		
7301	GGAGGGCTTA CCTCCCGAAT		CCAGTGCTGC GGTCACGACG		
7351	GCTCACCGGC CGAGTGGCCG		TCAGCAATAA AGTCGTTATT		
7401	GAGCGCAGAA CTCGCGTCTT				
7451	TTGTTGCCGG AACAACGGCC				
7501			CCGTAGCACC	ACAGTGCGAG	
	ATGGCTTCAT TACCGAAGTA	TCAGCTCCGG AGTCGAGGCC	TTCCCAACGA AAGGGTTGCT	TCAAGGCGAG AGTTCCGCTC	TTACATGATC AATGTACTAG
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7601	CCCCATGTTG GGGGTACAAC			CTTCGGTCCT GAAGCCAGGA		
7651	TCAGAAGTAA AGTCTTCATT	-		TCATGGTTAT AGTACCAATA		
7701	CATAATTCTC GTATTAAGAG			AGATGCTTTT TCTACGAAAA		
7751	TGAGTACTCA ACTCATGAGT	-		GTGTATGCGG CACATACGCC		
7801		CCGCAGTTAT	GCCCTATTAT	CCGCGCCACA GGCGCGGTGT	ATCGTCTTGA	
7851 - 🖸	TTAAAAGTGC AATTTTCACG	TCATCATTGG	AAAACGTTCT		AACTCTCAAG TTGAGAGTTC	
	GATCTTACCG CTAGAATGGC			GTAACCCACT CATTGGGTGA		
79 <u>51</u> . ≟	ACTGATCTTC TGACTAGAAG	TCGTAGAAAA	TGAAAGTGGT	CGCAAAGACC	CACTCGTTTT	
8001			TTTTTTCCCT	TATTCCCGCT		
80 <u>51</u>	AACTTATGAG	TATGAGAAGG	AAAAAGTTAT	AATAACTTCG		
81 01	GTTATTGTCT	GTACTCGCCT	ATGTATAAAC	AATGTATTTA TTACATAAAT	GAAAAATAAA CTTTTTATTT	
8151	CAAATAGGGG	TTCCGCGCAC AAGGCGCGTG	ATTTC			

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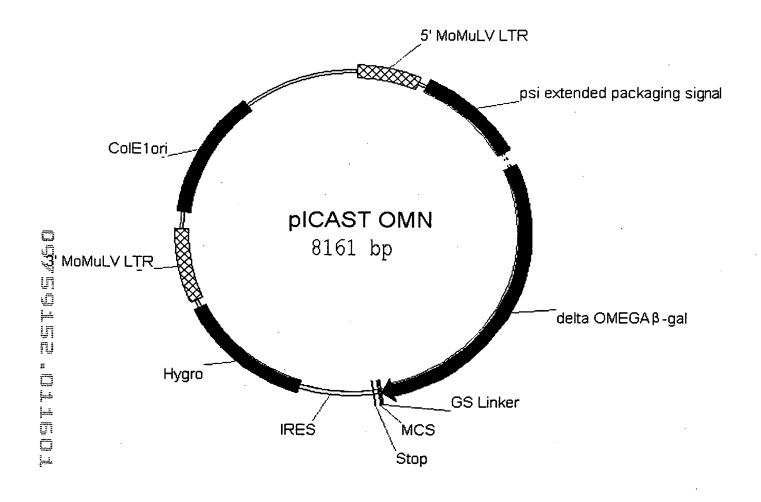


Figure 13A

l	CTGCAGCCTG GACGTCGGAC	AATATGGGCC TTATACCCGG				
51	CCCCGGCTCA GGGGCCGAGT	GGGCCAAGAA CCCGGTTCTT				
101	GGATATCTGT CCTATAGACA	GGTAAGCAGT CCATTCGTCA			TTCTTGTCTA	
151	GGTCCCCAGA CCAGGGGTCT	TGCGGTCCAG ACGCCAGGTC		-	ACCATCAGAT	
201	GTTTCCAGGG CAAAGGTCCC	TGCCCCAAGG ACGGGGTTCC				
251		GTTCGCTTCT CAAGCGAAGA				
3 0	GCTCAATAAA CGAGTTATTT					
14	TGACTGAGTC ACTGACTCAG					
401	CATCCGACTT GTAGGCTGAA	GTGGTCTCGC CACCAGAGCG				
451	TGACTACCCG ACTGATGGGC	TCAGCGGGGG AGTCGCCCCC				
5 01		CCCAGGGACC GGGTCCCTGG				
551	AGCAACTTAT	CTGTGTCTGT GACACAGACA				
601	TGCGCCTGCG ACGCGGACGC					
651	CGTGGTGGAA GCACCACCTT	CTGACGAGTT GACTGCTCAA			CTGGGAGACG	
701	TCCCAGGGAC AGGGTCCCTG	AAACCCCCGG	CAAAAACACC	GGGCTGGACT	CCTTCCCTCA	
751	CGATGTGGAA GCTACACCTT	TCCGACCCCG	TCAGGATATG AGTCCTATAC	TGGTTCTGGT ACCAAGACCA	AGGAGACGAG TCCTCTGCTC	
801	AACCTAAAAC TTGGATTTTG	AGTTCCCGCC TCAAGGGCGG	TCCGTCTGAA	TTTTTGCTTT	CGGTTTGGAA	
851	CCGAAGCCGC GGCTTCGGCG			AGCAAGACAC	AACAGAGACA	
901	CTGACTGTGT GACTGACACA	AAGACATAAA	CAGACTTTTA	ATCCCGGTCT	GACAATGGTG	

951	TCCCTTAAGT AGGGAATTCA			AGATGTCGAG TCTACAGCTC	
1001	ACAACCAGTC TGTTGGTCAG				CTTCTGCTCT GAAGACGAGA
1051	GCAGAATGGC CGTCTTACCG			CCGCGAGACG GGCGCTCTGC	
1101	CCGAGACCTC GGCTCTGGAG			GGTCTTTTCA CCAGAAAAGT	
1151	ATGGACACCC TACCTGTGGG			TGACCTGGGA ACTGGACCCT	
1201	TTTGACCCCC			GTACACCCTA CATGTGGGAT	
125基				CCTTGAACCT GGAACTTGGA	
	CCCCGCCTCG GGGGCGGAGC			TCACTCCTTC AGTGAGGAAG	
	GGCCGCTCTA CCGGCGAGAT		TGCTGAGTGA	TATCCCGCTA	
<u> </u>	TGCACCATCA ACGTGGTAGT		GTCGACGAAC	AGAAACTCAT	TTCCGAAGAA
	GACCTACTCG			TCACTGGCCG AGTGACCGGC	
1501	ACGTCGTGAC TGCAGCACTG			CCAACTTAAT GGTTGAATTA	
1551	CACATCCCC GTGTAGGGGG			GCGAAGAGGC CGCTTCTCCG	•
1601	CGCCCTTCCC GCGGGAAGGG			GGCGAATGGC CCGCTTACCG	
1651	GTTTCCGGCA CAAAGGCCGT			CTGGCTGGAG GACCGACCTC	
1701	CTGAGGCCGA GACTCCGGCT			ACTGGCAGAT TGACCGTCTA	
1751	GATGCGCCCA CTACGCGGGT			CCCATTACGG GGGTAATGCC	
1801	GTTTGTTCCC CAAACAAGGG			TTACTCGCTC AATGAGCGAG	
1851	TTGATGAAAG AACTACTTTC	GACCGATGTC	CTTCCGGTCT	GCGCTTAATA	

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1901	CAATTGAGCC		GTGGTGCAAC CACCACGTTG		
1951	CCAGGACAGT GGTCCTGTCA		CTGAATTTGA GACTTAAACT		
2001	CCGGAGAAAA GGCCTCTTTT		GTGATGGTGC CACTACCACG		
2051	TATCTGGAAG ATAGACCTTC		GTGGCGGATG CACCGCCTAC		
2101	CTCGTTGCTG GAGCAACGAC		CTACACAAAT GATGTGTTTA		
	CTCGCTTTAA GAGCGAAATT		AGCCGCGCTG TCGGCGCGAC		
2291	ATGTGCGGCG TACACGCCGC		CTACCTACGG GATGGATGCC		
2251			GCGGCACCGC CGCCGTGGCG		
2301			GCCGATCGCG CGGCTAGCGC		
2351	CTTTTGGGCT		CGCCGAAATC GCGGCTTTAG		
2401	GGTTGAACTG CCAACTTGAC	GTGTGGCGGC	ACGGCACGCT TGCCGTGCGA	CTAACTTCGT	
2451		GGCGCTCCAC	CGGATTGAAA GCCTAACTTT		
2501	GGCAAGCCGT CCGTTCGGCA		AGGCGTTAAC TCCGCAATTG		
2551	GCATGGTCAG CGTACCAGTC		AGCAGACGAT TCGTCTGCTA		
2601	TGAAGCAGAA ACTTCGTCTT		GCCGTGCGCT CGGCACGCGA		
2651	CCGCTGTGGT GGCGACACCA		CGACCGCTAC GCTGGCGATG		•
2701	AGCCAATATT TCGGTTATAA		GCATGGTGCC CGTACCACGG		
	ATCCGCGCTG TAGGCGCGAC	CGATGGCCGC	TACTCGCTTG	CGCATTGCGC	
	CGCGATCGTA GCGCTAGCAT	ATCACCCGAG TAGTGGGCTC		TGGTCGCTGG ACCAGCGACC	GGAATGAATC

2851	AGGCCACGGC TCCGGTGCCG	GCTAATCACG CGATTAGTGC				·
2901		CCCGGTGCAG GGGCCACGTC	ATACTTCCGC	CGCCTCGGCT	GTGGTGCCGG	
2951	ACCGATATTA TGGCTATAAT	TTTGCCCGAT AAACGGGCTA				
3001	CCCGGCTGTG GGGCCGACAC	CCGAAATGGT GGCTTTACCA				
3051	AGACGCGCCC TCTGCGCGGG	GCTGATCCTT CGACTAGGAA				
3101	CTTGGCGGTT GAACCGCCAA	TCGCTAAATA AGCGATTTAT				
31 <u>5</u> 1		TTCGTCTGGG				
- <u> </u>	TGTCCCGCCG	AAGCAGACCC	TGACCCACCT	AGTCAGCGAC	TAATTTATAC	
3201 	ATGAAAACGG TACTTTTGCC	CAACCCGTGG GTTGGGCACC			ACCGCTATGC	·
3251 TU	CCGAACGATC GGCTTGCTAG	GCCAGTTCTG CGGTCAAGAC				
-8						
3301	GCCGCATCCA CGGCGTAGGT	CGCGACTGCC				
- -		CCCCCNNACC		CCACCCAAMA	COMCOMMOCOCO	
3351		CGGGCAAACC GCCCGTTTGG				
3401	CATAGCGATA	ACGAGCTCCT	GCACTGGATG	GTGGCGCTGG	ATGGTAAGCC	
	GTATCGCTAT	TGCTCGAGGA	CGTGACCTAC	CACCGCGACC	TACCATTCGG	
3451	GCTGGCAAGC	GGTGAAGTGC	CTCTGGATGT	CGCTCCACAA	GGTAAACAGT	
	CGACCGTTCG	CCACTTCACG	GAGACCTACA	GCGAGGTGTT	CCATTTGTCA	
3501	TGATTGAACT	GCCTGAACTA	CCGCAGCCGG	AGAGCGCCGG	GCAACTCTGG	
		CGGACTTGAT				
3551	CTCACAGTAC	GCGTAGTGCA	ACCGAACGCG	ACCGCATGGT	CAGAAGCCGG	
	GAGTGTCATG	CGCATCACGT	TGGCTTGCGC	TGGCGTACCA	GTCTTCGGCC	* *
	GCACATCAGC					
	CGTGTAGTCG	CGGACCGTCG	TCACCGCAGA	CCGCCTTTTG		
	CGCTCCCCGC					
		GCGCAGGGTG		TAGACTGGTG	GTCGCTTTAC	
3701	GATTTTTGCA	TCGAGCTGGG	TAATAAGCGT			
. –		AGCTCGACCC				
3751	AGGCTTTCTT TCCGAAAGAA	TCACAGATGT AGTGTCTACA				

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3801			CGTGTCGATA GCACAGCTAT			
3851			CTTAATTAAC GAA'ITAATTG			
3901			TGCATTTCGA ACGTAAAGCT			
3951			GCCGTCTTTT CGGCAGAAAA			
4001			CGAGCATTCC GCTCGTAAGG			
4051			TTGAATGTCG AACTTACAGC			
4105			AACGTCTGTA TTGCAGACAT		CGTCCGTCGC	
4154			GGTGCCTCTG CCACGGAGAC			
4201			GCACAACCCC CGTGTTGGGG			
			ATGGCTCTCC TACCGAGAGG			·
43 - C	· · ·		TACCCCATTG ATGGGGTAAC			
4351			TGTGTTTAGT ACACAAATCA			
4401			GTGGTTTTCC CACCAAAAGG			
	ATGGTACTTT	TTCGGACTTG	TCACCGCGAC AGTGGCGCTG	CAGACAGCTC	TTCAAAGACT	
4501	AGCTTTTCAA	GCTGTCGCAG	TCCGACCTGA AGGCTGGACT	ACGTCGAGAG		
4551			CGATGTAGGA GCTACATCCT			
4601	CCATTTATCG	ACGCGGCTAC	GTTTCTACAA CAAAGATGTT	TCTAGCAATA	CAAATAGCCG	
4651	TGAAACGTAG	CCGGCGCGAG	CCGATTCCGG GGCTAAGGCC	TTCACGAACT	GTAACCCCTT	
4701	AAATCGCTCT	CGGACTGGAT	TTGCATCTCC AACGTAGAGG	GCGGCACGTG	TCCCACAGTG	

4751	GTTGCAAGAC CAACGTTCTG	CTGCCTGAAA GACGGACTTT	CCGAACTGCC GGCTTGACGG	CGCTGTTCTG GCGACAAGAC	CAGCCGGTCG GTCGGCCAGC	
4801	CGGAGGCCAT GCCTCCGGTA					
4851	TTCGGCCCAT AAGCCGGGTA	TCGGACCGCA AGCCTGGCGT	AGGAATCGGT TCCTTAGCCA	CAATACACTA GTTATGTGAT	CATGGCGTGA GTACCGCACT	
4901	TTTCATATGC AAAGTATACG	GCGATTGCTG CGCTAACGAC				
4951	TGGACGACAC ACCTGCTGTG	CGTCAGTGCG GCAGTCACGC	TCCGTCGCGC AGGCAGCGCG	AGGCTCTCGA TCCGAGAGCT	TGAGCTGATG ACTCGACTAC	
		AGGACTGCCC TCCTGACGGG	CGAAGTCCGG GCTTCAGGCC	CACCTCGTGC GTGGAGCACG	ACGCGGATTT TGCGCCTAAA	
505¶		AATGTCCTGA TTACAGGACT	CGGACAATGG GCCTGTTACC	CCGCATAACA GGCGTATTGT	GCGGTCATTG CGCCAGTAAC	
- 년 510隻 	ACTGGAGCGA	GGCGATGTTC CCGCTACAAG	GGGGATTCCC CCCCTAAGGG	AATACGAGGT TTATGCTCCA	CGCCAACATC GCGGTTGTAG	
5151		GGCCGTGGTT CCGGCACCAA			CGCGCTACTT GCGCGATGAA	
<u> </u>		CATCCGGAGC GTAGGCCTCG	TTGCAGGATC AACGTCCTAG	GCCGCGGCTC CGGCGCCGAG	CGGGCGTATA GCCCGCATAT	
52 5 1	TGCTCCGCAT ACGAGGCGTA				TGACGGCAAT ACTGCCGTTA	
- ‡#: 5301					TCGTCCGATC AGCAGGCTAG	
5351	CGGAGCCGGG GCCTCGGCCC	ACTGTCGGGC TGACAGCCCG	GTACACAAAT CATGTGTTTA	CGCCCGCAGA	AGCGCGGCCG TCGCGCCGGC	
5401					AAACCGACGC TTTGGCTGCG	
5451			TTTCCTTATC		GACCGGGATC	
5501			AATAAATCAG	AGGTCTTTTT	GGGGGGAATG	
5551	AAAGACCCCA TTTCTGGGGT				CGCCATTTTG GCGGTAAAAC	
5601	CAAGGCATGG GTTCCGTACC				AGATCAAGGT TCTAGTTCCA	
5651					ATCTGTGGTA TAGACACCAT	

5701	AGCAGTTCCT TCGTCAAGGA			ACAGATGGAA TGTCTACCTT		
5751	TGGGCCAAAC ACCCGGTTTG			TTCCTGCCCC AAGGACGGGG		
5801	CAAGAACAGA GTTCTTGTCT			GCCCTCAGCA CGGGAGTCGT		
5851	AACCATCAGA TTGGTAGTCT			GACCTGAAAT CTGGACTTTA		
5901	CTTATTTGAA GAATAAACTT			TCGCTTCTGT AGCGAAGACA		
5951	CTGCTCCCCG GACGAGGGGC			AACCCCTCAC TTGGGGAGTG		
- I -						
, e ^{rii}	AGTCCTCCGA TCAGGAGGCT			CCCGTGTATC GGGCACATAG		
6051				CTGTTCCTTG GACAAGGAAC		
- <u></u> 6101	CTCTGAGTGA GAGACTCACT			GTCTTTCATT CAGAAAGTAA		
_	GTATCAAAAT	TAATTTGGTT	TTTTTTCTTA	AGTATTTACA	TTAAATGGCC	
	CATAGITITA	ATTAAACCAA	AAAAAAGAAI	TCATAAATGT	AATTTACCGG	<u></u>
- <u>ļ</u> mi					COMMISSION	
62 <u>0</u> 1				GGGGAGAGGC CCCCTCTCCG		
6251				CTCGCTGCGC GAGCGACGCG		
					2 CCCERT TCC	
	GGCTGCGGCG CCGACGCCGC	TCGCCATAGT	CGAGTGAGTT	TCCGCCATTA		
6351		CCCTATTGCG	TCCTTTCTTG	ATGTGAGCAA TACACTCGTT	TTCCGGTCGT	
6401	AAAGGCCAGG	AACCGTAAAA	AGGCCGCGTT		TTCCATAGGC	
6451	TCCGCCCCCC AGGCGGGGG			GACGCTCAAG CTGCGAGTTC		
6501	CGAAACCCGA GCTTTGGGCT			GCGTTTCCCC CGCAAAGGGG		
6551	CCTCGTGCGC GGAGCACGCG		· ·	GCTTACCGGA CGAATGGCCT		
6601	CCTTTCTCCC GGAAAGAGGG			CTCATAGCTC GAGTATCGAG		

: : :

6651		CGGTGTAGGT GCCACATCCA				
6701		CAGCCCGACC GTCGGGCTGG	CGACGCGGAA			
6751	AGTCCAACCC TCAGGTTGGG	GGTAAGACAC CCATTCTGTG			TCGGTGACCA	
6801	AACAGGATTA TTGTCCTAAT	GCAGAGCGAG CGTCTCGCTC			AGTTCTTGAA TCAAGAACTT	
6851	GTGGTGGCCT CACCACCGGA	AACTACGGCT TTGATGCCGA		TTGTCATAAA	GGTATCTGCG CCATAGACGC	
6901	CTCTGCTGAA GAGACGACTT	GCCAGTTACC CGGTCAATGG		GAGTTGGTAG	CTCTTGATCC	
6951	GGCAAACAAA CCGTTTGTTT	CCACCGCTGG GGTGGCGACC				
1 52	GATTACGCGC CTAATGCGCG	TCTTTTTTC		TCTAGGAAAC	TAGAAAAGAT	
70 51	CGGGGTCTGA		AACGAAAACT	CACGTTAAGG	GATTTTGGTC	
7101		CAAAAAGGAT GTTTTTCCTA				
71 訂 - □ -		TATATATGAG ATATATACTC				
72 01	ATCAGTGAGG TAGTCACTCC	CACCTATCTC GTGGATAGAG				
7251	TGCCTGACTC ACGGACTGAG	CCCGTCGTGT GGGCAGCACA				
7301	CTGGCCCCAG GACCGGGGTC	TGCTGCAATG ACGACGTTAC				
7351		GTTATTTGGT	CGGTCGGCCT	TCCCGGCTCG	CGTCTTCACC	
7401		AATAGGCGGA	GGTAGGTCAG	ATAATTAACA	ACGGCCCTTC	
7451	CTAGAGTAAG GATCTCATTC	TAGTTCGCCA ATCAAGCGGT				
7501	GCTACAGGCA CGATGTCCGT	TCGTGGTGTC AGCACCACAG				
7551		GTTGCTAGTT	CCGCTCAATG	TACTAGGGGG	TACAACACGT	

· , · · ,

7601.	AAAAAGCGGT TTTTTCGCCA	TAGCTCCTTC ATCGAGGAAG			
7651	GCCGCAGTGT CGGCGTCACA	TATCACTCAT ATAGTGAGTA			
7701	TGTCATGCCA ACAGTACGGT	TCCGTAAGAT AGGCATTCTA			
7751	AGTCATTCTG TCAGTAAGAC	AGAATAGTGT TCTTATCACA		•	
7801	TCAATACGGG AGTTATGCCC	ATAATACCGC TATTATGGCG			
7851	CATTGGAAAA GTAACCTTTT	CGTTCTTCGG GCAAGAAGCC			
7901		TTCGATGTAA AAGCTACATT			
79 51		TCACCAGCGT AGTGGTCGCA	AAGACCCACT		
80 01		AAGGGAATAA TTCCCTTATT			
8051		TCAATATTAT AGTTATAATA			
8101		TATTTGAATG ATAAACTTAC			
81 51	GCGCACATTT CGCGTGTAAA	_			,

, :

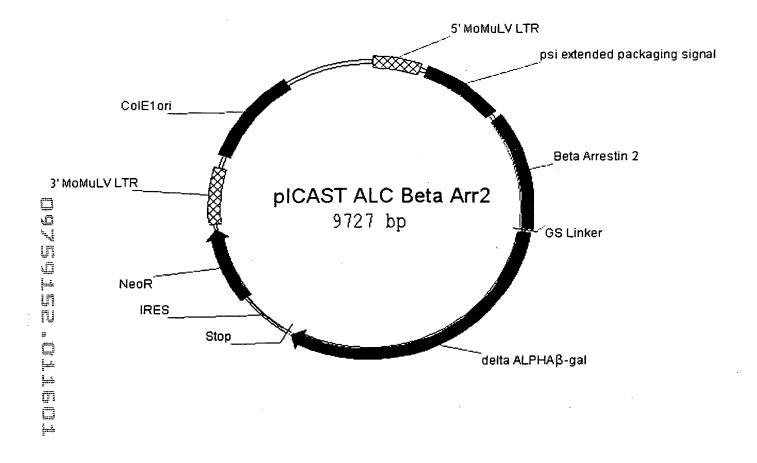


Figure 14

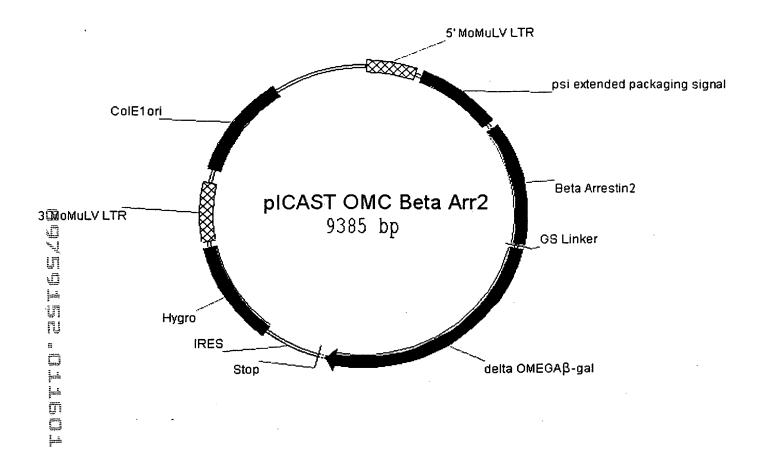


Figure 15

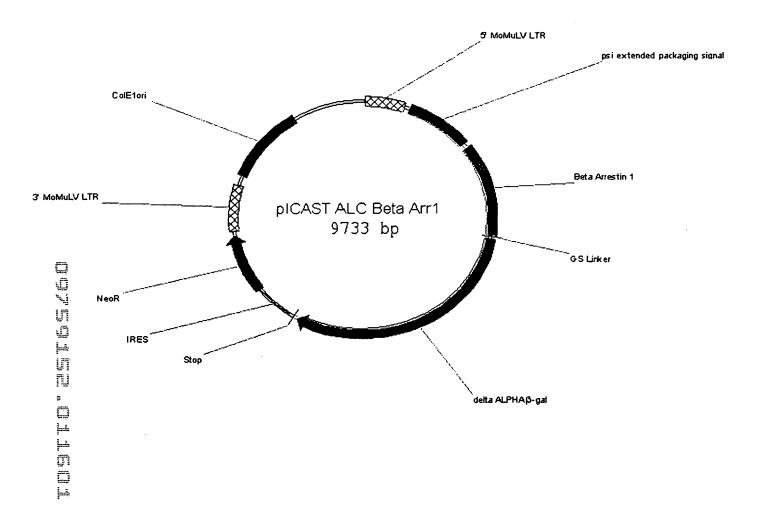


Figure 16

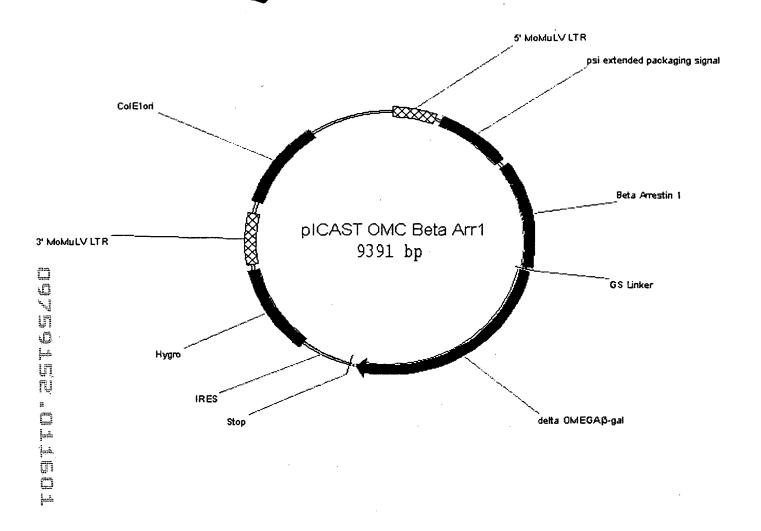


Figure 17

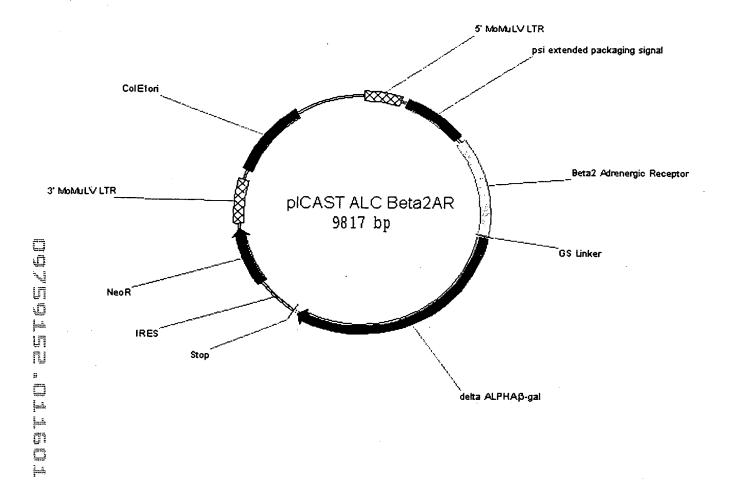


Figure 18

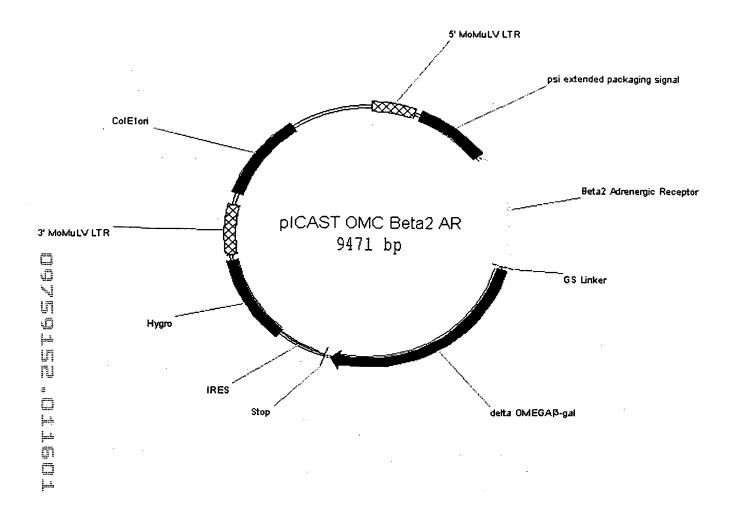


Figure 19

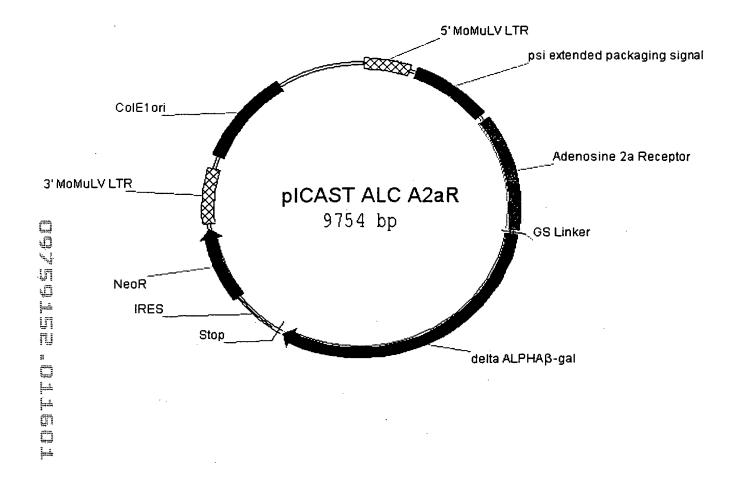


Figure 20

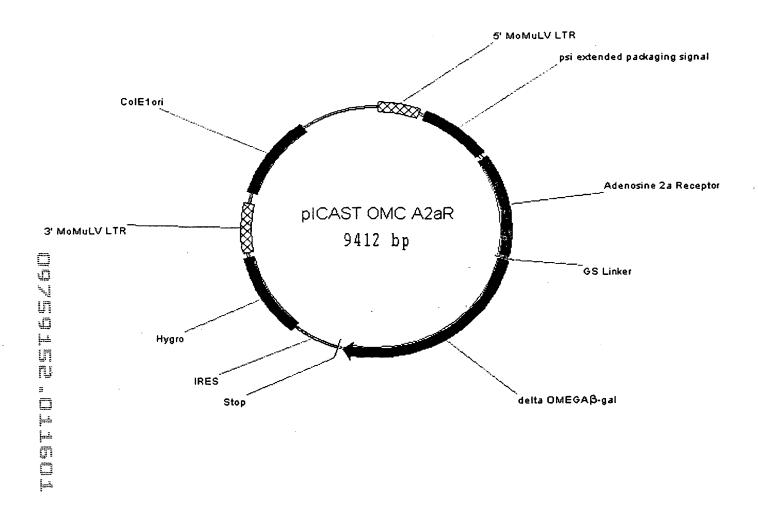


Figure 21

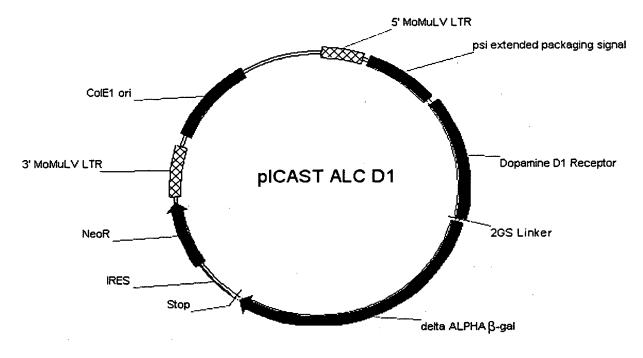


Figure 22

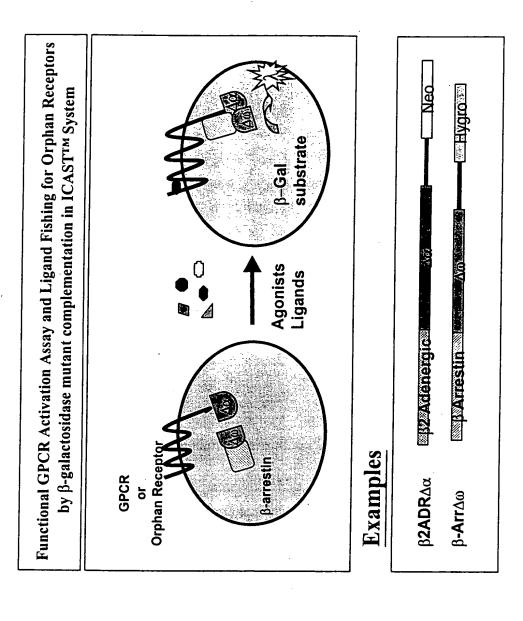
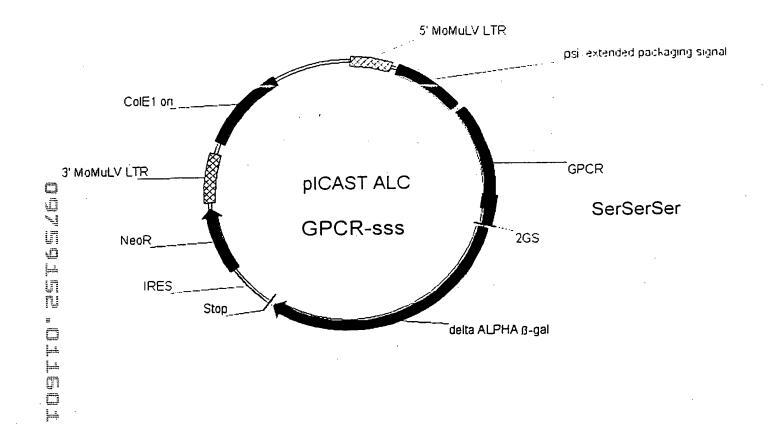
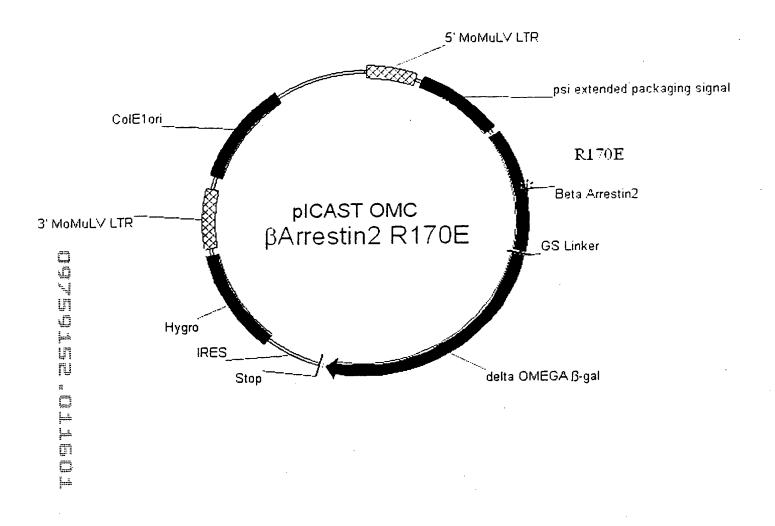


Figure 23



Vector for Expression of a GPCR with inserted Seronine/Threonine amino acid sequences as a fusion with $\beta\text{-gal}\ \Delta\alpha.$



Vector for Expression of mutant (R170E) $\beta\text{-arrestin2}\;$ as a fusion with $\beta\text{-gal}\;\Delta\omega.$

FIGURE 25

Phosphorylation Insensitive Mutant R170E $\beta\textsc{-Arrestin2}\Delta\omega$ Binds to $\beta2$ AR $\Delta\alpha$ in Response to Agonist Activation

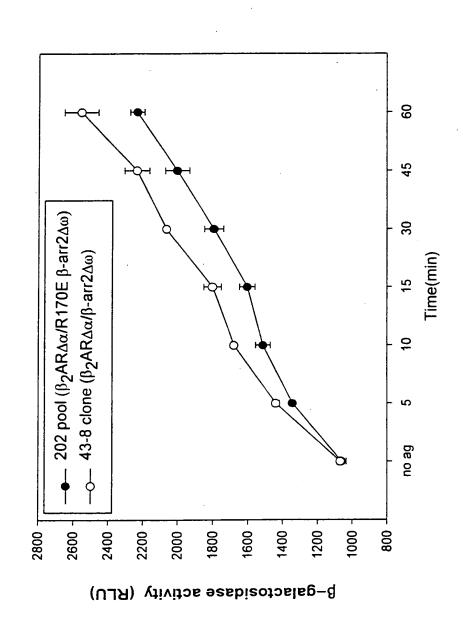
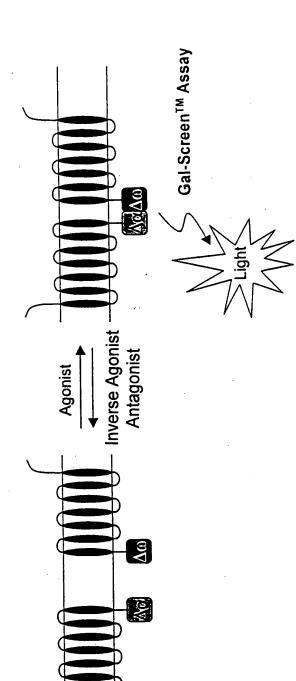
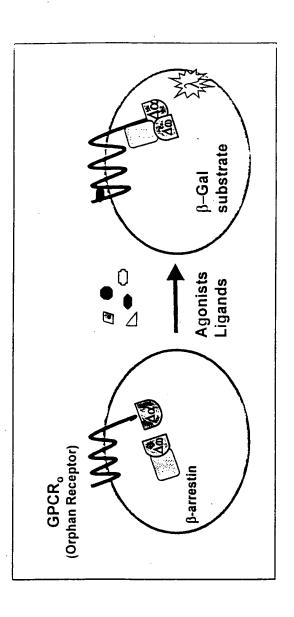


FIGURE 26



GPCR dimerization measured by β-gal complementation

FIGURE 27





Ligand Fishing for Orphan Receptors by β -galactosidase mutant complementation in ICAST^{FM} System

FIGURE 28